GenCore version 5.1.6 Copyright (c) 1993 - 2005 Compus

Compugen Ltd.

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                              Minimum DB seq length: Maximum DB seq length:
                                                                                                                                                                                                                                                                                                                   Total number of hits satisfying chosen parameters:
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                                                                                                                                                                                                                                                                                                                                                     2443163 segs, 439378781 residues
                                                                                                                                                                                                                                                                                                                                                                                      Gapop 10.0 , Gapext 0.5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-10-632-706-86
geneseqD2001s:*
geneseqD2002s:*
geneseqD2003as:*
geneseqD2001bs:*
geneseqD2004s:*
geneseqD2005s:*
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geneseqp2000s:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

24	23	22	21	20	19	18	17	16	15	14	13	12	11	10	9	8	7	o	5	4	ω	Ŋ	1	Result
456.5	456.5	456.5	456.5	456.5	456.5	456.5	456.5	456.5	456.5	460.5	460.5	•	460.5	460.5	460.5	٠	460.5	460.5		460.5	461.5	•	590	Score
77.4	77.4	77.4	77.4	77.4	77.4	77.4	77.4	77.4	77.4	78.1	78.1	78.1	78.1	78.1	78.1	78.1	78.1	78.1	78.1	78.1	78.2	86.5	100.0	Query Match Length
238	238	125	125	125	109	106	106	106	106	330	240	240	240	240	108	108	108	108	108	108	108	107	112	
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Aae25963 KDR bindi	Aae25961 KDR bindi	Mous	Chime		Anti	7	VEGF E	Anti				_	9 Antic	_		٠.	Antig	ഗ	_	Aay97236 Variable			Mouse 1	Description

45	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30	29	28	27	26	V.
439.5	439.5	443.5	443.5	445.5	447.5	447.5	447.5	447.5	447.5	447.5	448.5	448.5	448.5	449.5	449.5	449.5	453.5	456.5	456.5	400.5
74.5	74.5	75.2	75.2	75.5	75.8	75.8	75.8	75.8	75.8	75.8	76.0	76.0	76.0	76.2	76.2	76.2	76.9	77.4	77.4	//.4
109	105	236	106	110	129	129	129	108	107	107	258	256	108	266	235	108	125	238	238	200
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Aau81278	Aaw03182	Aay34096	Adr15137	Aar21305	Aay28357	Aay28385	Aar53329	Aar21290	Adz52586	Aab83159	Aab82898	VARPRIARY.	Aar21294	Adn / /30 /	Aaw41392	Aaw4 1 3 8 8	ABEZEST/	ACKIB222	AD] 26/32	
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## ALIGNMENTS

## RESULT 1 ADR38684 ADR38684 standard; peptide; 112 AA. ADR38684;

02-DEC-2004

(first entry)

antibacterial; antibody; botulinum neurotoxin type A; BONT/A; BONT/A neutralising epitope; anti-botulinum neurotoxin antibody; toxin neutralisation; botulinum neurotoxin poisoning; mouse; light chain variable region; single chain antibody; scFv. Mouse light chain variable region scFv seqid 86

Mus sp 09-SEP-2004. US2004175385-A1.

31-AUG-1998; 98US-00144886. 01-AUG-2002; 2002US-0400721P.

01-AUG-2003; 2003US-00632706

Marks JD, Amersdorfer P;

(REGC ) UNIV CALIFORNIA.

• • •

WPI; 2004-652009/63.

New isolated antibody that neutralizes botulinum neurotoxin type A, useful for diagnosing botulism or for treating pathologies associated with botulinum neurotoxin poisoning.

Example 1; SEQ ID NO 86; 110pp; English.

The invention describes an isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a specific clone where (I) binds to and neutralises botulinum neurotoxin type A (BoNT/A). An isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a clone chosen from clone 25, C25, C39, 1C6, 3D12, B4, 1F3, huC25, A1, Ar2, WR1(V), WR1(T), 3-1, 3-8, 3-10 and ING1, where (I) binds to and neutralizes botulinum neurotoxin type A (BoNT/A). Also described are: a polypeptide (II) comprising BoNT/A neutralising epitope having an epitope that is

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RESULT 2
ADR38683
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 100.0%; Score 590; DB 8; Best Local Similarity 100.0%; Pred. No. 1.6e-38; Matches 112; Conservative 0; Mismatches 0;
                                                                                                               New isolated antibody that neutralizes botulinum neurotoxin type A, useful for diagnosing botulism or for treating pathologies associated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     antibacterial; antibody; botulinum neurotoxin type A; BONT/A; BONT/A neutralising epitope; anti-botulinum neurotoxin antibody; toxin neutralisation; botulinum neurotoxin poisoning; mouse; light chain variable region; single chain antibody; scFv.
                                                       Example 1; SEQ ID NO 85; 110pp; English
                                                                                                                                                                         WPI; 2004-652009/63.
                                                                                                                                                                                                               Marks JD,
                                                                                                                                                                                                                                                                                          31-AUG-1998; 98US-00144886.
01-AUG-2002; 2002US-0400721P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADR38683 standard; peptide; 107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 112 AA,
                                                                                                                                                                                                                                                    (REGC ) UNIV CALIFORNIA.
                                                                                                 botulinum neurotoxin
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                                                                                                                                                                                                               Amersdorfer P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                               poisoning.
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The invention describes an isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a specific

Region

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CDR3 Framework CDR2

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Framework region

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Best Local S
Matches 101
                                                                                                                                                                                                                                                                                                                                                                                                               Single chain antibody; antibody; epidermal growth factor receptor; EGFR, tumour; cancer; glioma; melanoma; carcinoma; diagnosis; assessment; phage antibody library.
                                                                                                                                                                                  Region
                                                                                                                                                                                                                                                                    Region
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                                                                                                                                                                                                                                                                                                                                                                               Mus musculus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Anti-EGFR antibody light chain variable region (Clone L2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              protein; 108 AA
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91.0%;
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Pred. No. 2.4e-32;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Kettleborough AC, Bendig MM, Mitjans F, Rosell E, Blasco
                                                                                                                                                                                                                                                                                                                                                                                                  Immunoglobulin; antibody; complementary determing region; CDR; VEGF; vascular endothelial growth factor; KDR; vtascular endothelial growth factor; KDR; kinase insert domain containing receptor; multivalent; monovalent; kinase insert domain containing receptor; multivalent; monovalent; humanised antibody; chimeric antibody; tumour; diabody; triabody;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 108
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                                                                                                                                                                                                                                                                                                                                                                              glioblastoma multiforme; hemangioblastoma; AIDS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Variable light chain fragment of anti-SI(KDR) antibody.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19-DEC-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAY97236
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (MERE ) MERCK PATENT GMBH.
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02-DEC-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21-SEP-1995
  29-JAN-1999;
29-JAN-1999;
                                                                                                                                                                                          WO200044777-A1
                                                                                                                                                                                                                                                                      Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                  central nervous
                                                                                 28-JAN-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DIELTQSPAIMSASPGEKVTITCSASSSV-----SYMHWFQQKPGTSPKLWIYSTSNLAS
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                                                                                 2000WO-US002180
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94EP-00118970.
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  99US-00240736.
99US-0117726P.
                                                                                                                                                                                                                                                                                                                   system neoplasms; AIDS associated Karposi's sarcoma; deficiency syndrome; AIDS: human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          protein; 108 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            78.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 461.5; I
Pred. No. 1.6e
4; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 .6e-28;
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New immunoglobulin molecules are described that bind kinase insert domain containing receptor (KDR) with a comparable affinity to human vascular endothelial growth factor (VEGF). The antibodies neutralise KDR activation. The immunoglobulin may be a multivalent single chain antibody, a monovalent single chain antibody, a diabody, a triabody, a humanised antibody or a chimerised antibody. The immunoglobulin molecules bind specifically to an extracellular domain of the KDR receptor with the same affinity as VEGF. Overexpression of VEGF has been implicated in a number of human tumour cell lines including glioblastcoma multiforme, hemangish sarcoma. The antibodies therefore have applications in treating these conditions. This sequence encodes a preferred heavy chain complementary determining region of the immunoglobulins of the invention
                                                                                  Key
Region
                                                                                                                                                                                                            IMC-1C11; chimeric antibody; mouse; humān; antagonist; VEGF; vascular endothelial growth factor; angiogenesis; antiangioge: antitumour; neoplasm; collagen-vascular disease; autoimmune ditumour; breast carcinoma; lung carcinoma; prostate carcinoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 108 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 4; Page 51; 55pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel immunoglobulin molecules binding kinase insert domain-containing receptor with the same affinity as vascular endothelial growth factor,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI;
 Region
                                          Region
                                                                                                                                                                                                                                                                                   VEGF antagonist antibody IMC-1C11 light chain variable region
                                                                                                                                                                                                                                                                                                               11-SEP-2003
15-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                      AAB82710
                                                                                                                             Chimeric.
                                                                                                                                                       Mus sp.
                                                                                                                                                                                   glioblastoma
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        to reduce tumor growth.
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                                                                                                                                           sapiens.
                                                                                                                                                                                               carcinoma; ovarian
                                                                                                                                                                                                                                                                                                                                                                                                                                                            56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     61 GVPARFSGSGSGTSYSLTISRMEAEDAATYYCQQSNEDPYTFGSGDQAGNKR
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                                                                                                                                                                                                                                                                                                                                                                                    standard; protein;
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                                                                                                                                                                                   oma; ovarian carcinoma; neuroblastoma; multiforme; melanoma; therapy; light
                                                                                                                                                                                                                                                                                                               (revised)
(first entry)
                                          /label= CDR-L1
/note= "complementarity
49. .55
                                                                                               Location/Qualifiers
                note=
                             'label= CDR-L2
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82.1%;
                "complementarity
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Pred. No. 1.9e-28;
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                   determining region
                                                          determining
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                                                                                                                                                                                     light chain
                                                                                                                                                                                                                                          antiangiogenic;
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                                                          region
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                                                                                                                                                                                                                                 disease;
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RESULT 6
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(1011, a mouse-human chimeric antibody that has vascular endothelial
(1011) growth factor (VEGP) antagonist activity. The antibody, or a fragment of
(1011) the used as an anti-angiogenic molecule, together with a
(1011) condition in a mammal, especially a human. The invention relates
(1012) condition in a mammal, especially a human. The invention relates
(1013) condition by administering an anti-angiogenic molecule and a
(1014) condition by administering an anti-angiogenic molecule and a
(1014) condition while minimising or preventing significant toxicity of the
(1015) condition while minimising or preventing significant toxicity of the
(1015) condition while minimising or preventing significant toxicity of the
(1016) condition while minimising or preventing significant toxicity of the
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(1016) condition while minimising or 
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2003 to standardise OS field)
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Mouse; immunoglobulin; kinase insert domain-containing receptor; KDR;
                                                Mouse anti-KDR plCl1 scFv antibody VL region #1
                                                                                                   15-NOV-2002
                                                                                                                                                                                                  AAE25956 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 108 AA;
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31-MAR-2000; 2000US-00539692.
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                                                                                                                                                                                                                                                                                                                                                    GVPARFSGSGSGTSYSLTISRMEAEDAATYYCQQSNEDPYTFGSGDQAGNKR 112
                                                                                                                                                                                                                                                                                                                                                                                                                                                           DIELTQSPAIMSASPGEKVTTTCRASESVDSYGHSFMQWFQQKPGTSPKLWIYSTSNLAS
                                                                                                                                                                                                                                                                                                                      GVPARFSGSGSGTSYSLTISRMEABDAATYYCOORSSYPFTFGSGTKLEIKR 107
                                                                                                                                                                                                                                                                                                                                                                                                                          DIELTQSPAIMSASPGEKVTITCSASSSV----SYMHWFQQKPGTSPKLWIYSTSNLAS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                 (first entry)
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                                                                                                                                                                                                protein; 108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             78.1%;
82.1%;
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Pred. No. 1.9e-28;
4; Mismatches 11
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AAU74413
ID AAU74413
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention relates to novel immunoglobulin molecules that bind to kinase insert domain-containing receptor (KDR) (a human homologue of mouse foetal liver kinase (FLK)-1 receptor) with an affinity comparable to human vascular endothelial growth factor (VEGF) and that neutralises activation of KDR. Sequences of the invention are useful for neutralising the activation of KDR, for reducing tumour growth and for inhibiting angiogenesis. The present sequence is mouse anti-KDR plCil scFv antibody light chain variable region (VL)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel immunoglobulin molecule for reducifig tumor growth, binds to kinase insert domain-containing receptor with an affinity comparable to human vascular endothelial growth factor, and fieutralizes activation of KDR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   foetal liver kinase; FLK-1 receptor; vastular endothelial growth factor;
VEGF; tumour growth; light chain variable region; VL; angiogenesis;
plCl1; scFv antibody.
                                                                                                                           Antigen-binding protein; antibody light chain variable cytostatic; angiogenesis inhibitor; tumour; leukaemia; vascular endothelial growth factor receptor; VEGF;
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N-PSDB; AAD42821.
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                                                                                                                                                                                                                                                                                         AAU74413 standard; peptide; 108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 108 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 4; Page 11; 34pp; English.
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                                                                                                                                                                                         Antigen-binding
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local
                                                                                                              proliferation inhibitor
                                                                                                                                                                                                                                                                                                                                                                         56 GVPARFSGSGSGTSYSLTISRMEAEDAATYYCQQRSSYPFTFGSGTKLEIKR 107
                                                                                                                                                                                                                                                                                                                                                                                                    61 GVPARFSGSGSGTSYSLTISRMEAEDAATYYCOOSNEDPYTFGSGDQAGNKR 112
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ) ZHU Z.
) WITTE L.
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                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                           protein light chain variable domain (VH)
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82.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 460.5; DB 5;
Pred. No. 1,9e-28;
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WO200190192-A2.

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RESULT 8
ABJ26725
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Crimplex of two polypeptides (P1) and two second polypeptides (P2) which care stably associated in an immunoglobulin like complex. P1 has an CC antigen-binding site located to the N terminus of immunoglobulin (Ig) (Ig) chain constant domain (CL domain), and P2 has an antigen-binding CC light chain constant domain (CL domain), and P2 has an antigen-binding CC site located to the N terminus of the CH1 domain. (I) is useful for: CC neutralising the activation of a vascular endothelial growth factor CC (VEGP) receptor; reducing tumour growth; inhibiting angiogenesis; creducing endothelial cell proliferation; inhibiting VEGP induced CC migration of human leukaemia cells; blocking interaction of a protein and it sligand; promoting interactions between immune cells and target cells; cand in vivo and in vitro for investigative, diagnostic or treatment CC methods. The design of (I) provides for efficient production so that CC substantially all of the antigen-binding proteins produced are assembled in the desired configuration. (I) is bivalent and bispecific, homogeneous and in tetrameric form. The heavy chain constant domains which constitute the Fc region (e.g., CH2 and CH3 for an IGG molecule) of a natural constitute of an engineered provide other antibody functions can be present. There is no requirement for processing in vitro to obtain the complete product. This sequence represents a light chain variable domain (VH) incorporated contor Fv, an engineered processing in vitro to obtain the complete product. The method of the invention
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New bispecific immunoglobulin-like antigen-binding protein for reducing tumor growth and for inhibiting angiogenesis, comprises a complex of two polypeptides and two second polypeptides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   24-MAY-2000; 2000US-0206749P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 108 AA;
                                                                      Cytostatic; antibody; antigen binding site; VEGF receptor; mitogenesis; leukaemia cell; vascular endothelial growth factor; tumour; bispecific antigen-binding protein; mouse; murine.
                                  Mus sp
                                                                    bispecific
                                                                                                                                            VEGF binding
                                                                                                                                                                                01-MAY-2003
                                                                                                                                                                                                                                                    ABJ26725 standard; protein; 108 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              57; Page 57;
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                                                                                                                                                                                                                                                                                                                                                                | GVPARFSGSGSGTSYSLTISRMEAEDAATYYCQQSNEDPYTFGSGDQAGNKR 112
                                                                                                                                                                                                                                                                                                                                             GVPARPSGSGSGTSYSLTISRMEAEDAATYYCQQRSSYPFTFGSGTKLEIKR 107
                                                                                                                                                                                                                                                                                                                                                                                                                   DIELTQSPAIMSASPGEKVTITCSASSSV----SYMHWFQQKPGTSPKLWIYSTSNLAS
                                                                                                                                                                                                                                                                                                                                                                                                                                        DIELTQSPAIMSASPGEKVTTTCRASESVDSYGHSFMQWFQQKPGTSPKLMIYSTSNLAS
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82.1%;
                                                                                                                                            protein
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                                                                    protein;
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Pred. No. 1.9e
4; Mismatches
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                                                                                                                                              SEQ ID
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د.
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WO2003002144-A1

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RESULT 9
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RESULT 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 tumours and for in vivo or in vitro for investigative and diagnostic methods. This sequence represents a mouse protein relating to the bispecific antibodies that bind to the VEGF receptors of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New bispecific antibodies having antigen-binding sites specific for a first vascular endothelial growth factor (VEGF) receptor and for a second VEGF receptor, useful for inhibiting migration of leukemia cells, or for treating tumors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to a novel antibody having a first antigen binding site specific for a first vascular endothelial growth factor (VEGF)
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                                                                                                                                                                                                                                                                                                                                                                           antibody; antigen binding site; vascular endothelial growth factor; VEGF receptor; immunoglobulin; tumour; anglogenesis; cytostatic; immunostimulant; vaccine; complementarity determining region; variable;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADK18202 standard; peptide; 108 AA.
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                                                         26-JUN-2002; 2002WO-US020332
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mouse VEGF
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                                                                                                                      24-DEC-2002;
      (IMCL-) IMCLONE SYSTEMS INC
                                                                                                                                                                                                                                                                                                                                                      light;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 92;
                                                                                                                                                                                                                                                                                                                                                         mouse; murine.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DIELTQSPAIMSASPGEKVTITCSASSSV-----SYMHWFQQKPGTSPKLWIYSTSNLAS 55
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  receptor light chain variable domain
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82.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 460.5; DB 6;
Pred. No. 1.9e-28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        peptide,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length
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RESULT 10
AAE25960
ID AAE25
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AC AAE25
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IS-NO
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MOUSE
KW MOUSE
KW FOCTS
KW VEGF;
KW VEGF;
KW VEGF;
KW VEGF
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Best Local (
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                                                                                                                                                                                    Mouse; immunoglobulin; kinase insert domain-containing receptor; KD foetal liver kinase; FLK-1 receptor; vascular endothelial growth fa VEGF; tumour growth; heavy chain variable region; VH; angiogenesis;
                                                                                                                                                                                                                                                                                  KDR binding
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                          30-MAY-2002
                                                                      US2002064528-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                        AAE25960 standard; protein; 240 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 108
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                                                                                                                                                                ; scFv a
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                                                                                                                                                                   antibody
                                                                                                                                                                                                                                                                                  immunoglobulin related mouse
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82.1%;
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Pred. No. 1.9
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                                                                                                                                                                                                                                                                                  protein
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ARESULT 11
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Best Local
                                                                                                                                                                                                                                            Antigen-binding protein; single chain variable fragment; scFv; cytostatic; angiogenesis inhibitor; tumour; leukaemia; antibody vascular endothelial growth factor receptor; VBGF; cell proliferation inhibitor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention relates to novel immunoglobulin molecules that bind to kinase insert domain-containing receptor (KDR) (a human homologue of mouse foetal liver kinase (FLK)-1 receptor) with an affinity comparable to human vascular endothelial growth factor (VEGF) and that neutralises activation of KDR. Sequences of the invention are useful for neutralising the activation of KDR, for reducing tumour growth and for inhibiting angiogenesis. The present sequence is KDR binding immunoglobulin related
                                                                                                                                                                                                      Mus sp.
Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 16-17; 34pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel immunoglobulin molecule for reducing tumor growth, binds to kinase insert domain-containing receptor with an affinity comparable to human vascular endothelial growth factor, and neutralizes activation of KDR.
                                                                                                   Region
                                                                                                                                                                                                                                                                                                                      Antigen-binding protein, single chain variable fragment version #1.
                                                                                                                                                                                                                                                                                                                                                                                 AAU74419;
                                                                                                                                                                                                                                                                                                                                                                                                            AAU74419 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 240 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DIELTQSPAIMSASPGEKVTITCSASSSV----SYMHWFQQKPGTSPKLWIYSTSNLAS
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                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                     /note= "15 amino a of the single chai AAS20285"
                                                                                                                                                                         Location/Qualifiers
                                                                                                                 /note= "Heavy
in claim 57"
note= "Light chain variable domain. Specifically claimed
                                                                                    /label= Linker
                                                                                                                                            'label=
                                                                                                                                                                                                                                                                                                                                                                                                            protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 78.1%;
82.1%;
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Pred. No. 4.2e-28;
                                                         chain
                                                                                                                               chain variable domain. Specifically claimed
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                                                       acid linker
sin variable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   11;
                                                       joins the VH and VL regions fragment protein. Encoded by
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د.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    239
                                                                                                                                                                                                                                                                                            antigen;
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RESULT 12
ABJ26731
ID ABJ26
XX ABJ26
XX ABJ26
XX UEGF
XX VEGF

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CC antigen-binding site located to the N terminus of immunoglobulin (Ig)
CC light chain constant domain (CL domain), and P2 has an antigen-binding
CC site located to the N terminus of the CHI domain. (I) is useful for:
CC enutralising the activation of a vascular endothelial growth factor
CC (VEGP) receptor; reducing tumour growth; inhibiting angiogenesis;
CC reducing endothelial cell proliferation; inhibiting angiogenesis;
CC enducing endothelial cell proliferation; inhibiting vEGF induced
CC its ligand; promoting interactions between immune cells and target cells;
CC and in vivo and in vitro for investigative, diagnostic or treatment
CC enthods. The design of (I) provides for efficient production so that
CC in the desired configuration. (I) is bivalent and bispecific, homogeneous
CC in the desired configuration. (I) is bivalent and bispecific, homogeneous
CC antibody and which provide other antibody functions can be present. There
CC is no requirement for processing in vitro to obtain the complete product.
CC is the samino acid sequence of a single chain variable fragment
CC (scPv), an engineered protein containing a variable light and variable
CC heavy domain on one polypeptide, described in the method of the invention
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                tumor growth and for inhibiting angiogene polypeptides and two second polypeptides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New bispecific immunoglobulin-like antigen-binding protein for reducing tumor growth and for inhibiting angiogenesis, comprises a complex of two
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2002-106189/14.
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Cytostatic; antibody; antigen binding site; VEGF receptor; mitogenesis; leukaemia cell; vascular endothelial growth factor; tumour; bispecific antigen-binding protein; mouse; murine.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 240 AA;
                                                                                                                                                                                                 01-MAY-2003
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                                                                                                                              VEGF binding related protein SEQ ID No
                                                                                                                                                                                                                                                                                                                            ABJ26731 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GVPARFSGSGSGTSYSLTISRMEAEDAATYYCQQSNEDPYTFGSGDQAGNKR 112
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DIELTOS PAIMSAS POEKVTITCS ASS SV-----SYMHWFQQXFGTS PXLWIYSTSNLAS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GVPARFSGSGSGTSYSLTISRMEAEDAATYYCQQRSSYPFTFGSGTKLEIKR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                              (first entry)
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                                                                                                                                                                                                                                                                                                                         protein; 240
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82.1%;
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Pred. No. 4.2e-28;
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                                                                                                                                     27.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 5;
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ADK1821
ID ADK1821
AC ADK18
AC ADK18
XX MOUSE
XX MOUSE
XX MOUSE
XX MOUSE
XX MOUSE
XX MOUSE
XX WOEPV,
XX WO200
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to a novel antibody having a first antigen binding site specific for a first vascular endothelial growth factor (VEGF) receptor and a second antigen-binding site specific for a second VEGF receptor. The bispecific antigen-binding proteins block activation of the VEGF receptor and are useful for reducing or inhibiting VEGF-induced cellular functions such as mitogenesis of vascular endothelial cells and migration of leukaemia cells. The antibodies are useful for treating tumours and for in vivo or in vitro for investigative and diagnostic methods. This sequence represents a mouse protein relating to the bispecific antibodies that bind to the VEGF receptors of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New bispecific antibodies having antigen-binding sites specific for a first vascular endothelial growth factor (VEGF) receptor and for a second VEGF receptor, useful for inhibiting migration of leukemia cells, or for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Zhu
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                                                                                                                                                           antibody; antigen binding site; vascular endothelial growth factor; VEGF receptor; immunoglobulin; tumour; angiogenesis; cytostatic; immunostimulant; vaccine; KDR; kinase insert domain-containing rece
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 240 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 55-56; 98pp; English
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                                                                                                             de snw
                                                                                                                                                                                                                           Mouse kinase insert domain-containing receptor scrv protein,
                                                                                                                                                                                                                                                           06-MAY-2004
                                                                                                                                                                                                                                                                                           ADK18221;
                                                                                                                                                                                                                                                                                                                         ADK18221 standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (IMCL-)
                                                                               WO2004003211-A1
                                                                                                                                           scrv; mouse;
                                                                                                                                           .mmunostimulant; vaccine; KDR; kinase
scFv; mouse; murine.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     133 DIBLTQSPAIMSASPGEKVTITCSASSSV----SYMHWFQQKPGTSPKLWIYSTSNLAS
                                                                                                                                                                                                                                                                                                                                                                                                          188
                                                                                                                                                                                                                                                                                                                                                                                                                        1 DIRLTQSPAIMSASPGEKVTTTCRASESVDSYGHSFMQWFQQKPGTSPKLWIYSTSNLAS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       92;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     tumors.
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                                                                                                                                                                                                                                                           (first
                                                                                                                                                                                                                                                            entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      78.1%;
82.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 460.5; DB 6
Pred. No. 4.2e-28;
4; Mismatches 11
                                                                                                                                                                                                                                                                                                                             240
                                                                                                                                                                                                                                                                                                                             8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          240;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          5;
                                                                                                                                                                                                                                SEQ ID
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24-DEC-2002; 2002WO-US041372.

08-JAN-2004.

26-JUN-2002; 2002WO-US020332

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RESULT 14
AAB70842
ID AAB70
XX AAB70
XX AAB70
XX I1-SE
DT 11-SE
DT 25-JU
XX T 1yn
KW Cell:
KW Gene
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to a novel antibody having a first antigen binding CC site specific for a first vascular endothalial growth factor (VEGF) CC receptor and a second antigen specific for a second VEGF receptor. The CC invention further relates to: an antibody that specifically binds to an extracellular domain of a first VEGF receptor and an extracellular domain CC of a second VEGF receptor, where binding of the antibody to the first and CC second VEGF receptor neutralizes activation of that VEGF receptor; a cC method for making the antibody by co-expressing in a host cell a cC immunoglobulin heavy chain domain located to the N-terminus of the second immunoglobulin light chain variable domain; a recombinant DNA construct CC encoding a second polypeptide having the second immunoglobulin light chain variable domain, a recombinant DNA construct CC immunoglobulin light chain variable domain, for time and in a manner CC sufficient to allow expression of the polypeptides and formation of the second VEGF receptor in a cell by treating a cell with the antibody; a method for neutralizing activation of a first VEGF receptor CC antibody; a method for recursive timour growth in a mammal by treating the mammal by treating a cell with the cCC antibody cited above; a method for reducing tumour growth in a mammal by treating a cell with the activities. The VEGF receptor antigen can be used to create a vaccine. The antibody cited above. The antibody has cytostatic and immunostimulant combining angiogenesis in a mammal by treating the mammal with the activities. The VEGF receptor antigen can be used to create a vaccine. The antibody is useful for reducing tumor growth in a mammal and for inhibiting angiogenesis. This secuence represents a kinase insert domain-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local S
Matches 92
                                                              cell-targeting vector; antiviral; cytostatic; immunostimulant; AIDS; gene therapy; immunization; diagnosis; T cell-associated disease; SC acquired immune deficiency syndrome; severe combined immune deficien
                                                                                                                                                                                  SNV-env leader/human 7B2-scFv fusion construct.
                                                                                                                                                                                                                                   11-SEP-2003
25-JUN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New antibody having a first and second binding sites specific for a first and second vascular endothelial growth factor (VEGF) receptor, useful for reducing tumor growth in a mammal and for inhibiting angiogenesis.
  Homo sapiens
                                                                                                                                     T lymphocyte; antibody; single chain variable antibody; scFv; human;
                                                                                                                                                                                                                                                                                                         AAB70842;
                                                                                                                                                                                                                                                                                                                                                       AAB70842 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 240 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            inhibiting anglogenesis. This sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; SEQ ID NO 27; 99pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      188
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            61 GVPARFSGSGSGTSYSLTISRMEAEDAATYYCQQSNEDPYTFGSGDQAGNKR 112
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 DIELTQSPAIMSASPGEKVTTTCRASESVDSYGHSFMQWFQQKPGTSPKLWIYSTSNLAS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       92; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GVPARFSGSGGTSYSLTISRMEAEDAATYYCQQRSSYPFTFGSGTKLEIKR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DIELTOS PAIMS AS PGEKVTITCS ASSSV-----SYMMWFOOKPGTS PKLWIYSTSNLAS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                         (revised)
                                                                                                                                                                                                                                                                                                                                                     protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          78.1%;
82.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This sequence represents a kinase insert domain-
protein of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 460.5; DB 8
Pred. No. 4.2e-28;
4; Mismatches 11
                                                                                                                                                                                                                                                                                                                                                       330
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     240;
                                                                      deficiency;
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                                                                                           SCID;
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11-SEP-2003

(revised)

AAE13144,

AAE13144 standard;

protein; 106 AA

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RESULT 15
AAE13144
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                                                                                                                                                                                                                                                                                                      This invention describes a novel cell-targeting vector (A) containing a CDNA sequence (I) encoding a single-chain variable antibody fragment (G (scFv). The products of the invention have antiviral, cytostatic and cimmunostimulant activity and can be used in gene therapy, immunization C and diagnosis particularly of T cell-associated diseases, specifically C acquired immune deficiency syndrome (AIDS), severe combined immune C confidency syndrome (AIDS), severe combined immune C confidency (SCID) or T cell lymphoma. (A) target T cells, independently C confidency (SCID) or T cell lymphoma. (A) target T cells, independently C confidency (SCID) or T cell lymphoma. (A) target T cells, independently C confidency (SCID) or T cell lymphoma. (A) target T cells, independently C confidency (SCID) or T cell lymphomatric or ther human cells. A vector designated 7A5 encodes a 329 amino acid single-chain variable antibody C confinency, fully defined in the specification. It was used to transform C confidency (SNV)) cells, C8166 (human T lymphocyte) cells, and Hela (human cervical carcinoma) cells. After 48 hours, the cells were stained with X-gal to C construction for D17, 1 million for C8166 but less than 100 for Hela, C showing the high selectivity for human T cells. This sequence represents the SNV-env leader/human 7B2-scFv fusion construct used in the invention. C (Updated on 11-SEP-2003 to standardise OS field)
                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                              Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     of e.g. acquired immune def variable antibody fragment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cell-targeting vector selective for T lymphocytes, useful in gene therapy of e.g. acquired immune deficiency syndrome, encodes a single-chain
                                                                                                                                                                                                                                                                                                    Sequence 330 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 1; Fig 3; 18pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cichutek K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       29-MAR-2001.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Key
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Spleen necrosis virus Chimeric.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             27-SEP-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Protein
                                                                                                                                                     181
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                                                                           236 GVPARFSGSGSGTSYSLTISRMEAEDAATYYCOORSSYPFTFGSGTKLEIKR
                                                                                                            61 GVPARFSGSGSGTSYSLTISRMEAEDAATYYCQQSNEDPYTFGSGDQAGNKR 112
                                                                                                                                                                          1 DIELTQSPAIMSASPGEKVTTTCRASESVDSYGHSFMQWFQQKPGTSPKLWIYSTSNLAS
                                                                                                                                                                                                                            92;
                                                                                                                                                                                                                                                Similarity
                                                                                                                                                   DIELTQSPAINSASPGEKVTITCSASSSV----SYMHWFQQKPGTSPKLWIYSTSNLAS
                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Engelstaedter M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             99DE-01046142
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /label= SNV-env_leader
                                                                                                                                                                                                                                            78.1%;
82.1%;
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                                                                                                                                                                                                                                              Score 460.5;
Pred. No. 5.8
                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                .8e-28;
                                                                                                                                                                                                                                                             DB 4;
                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                               Length 330;
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                                                                                                                                                                                                                            Gaps
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28-JAN-2002 (first entry)

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                                                                                                                                                                                                                                                                                                                                                             The invention relates to a method for inhibiting the growth of non-solid cc tumour cells that are stimulated by a ligand of vascular endothelial cc growth factor receptor (VEGFR) in mammals particularly humans. The method crown factor receptor (VEGFR) in mammals particularly humans. The method crown factor receptor (VEGFR) in mammals particularly humans. The method cc involves treating the mammals with humanised VEGFR monoclonal antibodies cc (antagonists). Humanised monoclonal antibody comprises humanised mouse controls region of the humanised constant region, where the humanised cc (CC variable region contains mouse complementarity determining region cc (CDR) grafted into human variable region. The method is useful for creating leukaemias such as acute or chronic myelocytic leukaemia, acute cor chronic lymphocytic leukaemia, erythrocytic or monocytic leukaemia, cc multiple myelomas and lymphoid cells, particularly those related to non-cc Hodgkin's and Hodgkin's disease. The present sequence is humanised cc antibody light chain fragment used in the exemplification of the crown to the control of the
                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Vascular endothelial growth factor receptor; VEGFR; antagonist; tumour; cytostatic; light chain; myelocytic leukaemia; lymphocytic; erythrocytic; monocytic; multiple myeloma; lymphoid cell; Hodgkin's disease; mouse; human; chimeric.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Inhibiting growth of non-solid tumor cells useful to treat bone marrow tumors such as leukemias or multiple myeloma comprises treatment with antagonist of a vascular endothelial growth factor receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2001-662942/76.
N-PSDB; AAD21670.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 8; Page 16; 68pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          31-MAR-2000; 2000US-00540770.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              30-MAR-2001; 2001WO-US010504.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   11-OCT-2001.
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                                                                                                                                                                                                                                                                                                                      Sequence 106 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Witte L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Chimeric.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (IMCL-) IMCLONE SYSTEMS INC. (CORR ) CORNELL RES FOUND INC.
                                                                                                                                                                                                                                        Match 77.4%;
Local Similarity 85.7%;
  56
                                         61 GVPARFSGSGSGTSYSLTISRMEAEDAATYYCQQSNEDPYTFGSG 105
                                                                                                   GVPARFSGSGSTSYSLTISRMEAEDAATYYCQQRSSYPFTFGSG 100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Rafii S;
                                                                                                                                                                                                                  Conservative
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                                                                                                                                                                                                             Score 456.5; DB 4; Length 106; Pred. No. 3.9e-28; 3; Mismatches 7; Indels 5
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Search completed: November 21, 2005, 12:19:54 Job time : 151.949 Becs

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Result
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Perfect score:
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Maximum DB seq length: 200000000
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 Pred. No. is the number of results predicted by chance to have a
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Gapop 10.0 , Gapext 0.5
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2: pir2:*
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 GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
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 BB
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S2636660
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       283416
 Description
g kappa chain V r
g kappa chain pre
g light chain V r
g kappa chain Pre
g kappa chain Pre
g kappa chain V r
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B54378
Ig light chain V region anti-triplex DNA - mouse (fragment)
C;Bpecies: Mus musculus (house mouse)
C;Date: 06-Oct-1994 #sequence_revision 18-Nov-1994 #text_change 21-Jan-2000
C;Accession: B54378
R;Agazie, Y.M.; Lee, J.S.; Burkholder, G.D.
J. Biol.: Chem. 269, 7019-7023, 1994
A;Title: Characterization of a new monoclonal antibody to triplex DNA and immunofluoresc A;Reference number: A54378; MUID:94165109; PMID:7509814
A;Reference number: A54378; MUID:94165109; PMID:7509814
A;Accession: B54378
A;Accession: B54378
A;Status: preliminary
A;Common A; Reference number: A54378; MUID:94165109; PMID:7509814
                                                                                                                                                                                                                      Ig kappa chain precursor V region (MRL22) - mouse C;Species: Mus musculus (house mouse) C;Date: 21-Man-1990 #sequence_revision 31-Dec-1990 #text_change 21-Jan-2000 C;Date: 21-Man-1990 #sequence_revision 31-Dec-1990 #text_change 21-Jan-2000 C;Accession: A3513 R;Kofler, R.; Strohal, R.; Balderas, R.S.; Johnson, M.E.; Noonan, D.J.; Duchosal, M.J. Clin. Invest. 82, 852-860, 1988 A;Title: Immunoglobulin kappa light chain variable region gene complex organization A;Reference number: A94689; MUID:88331394; PMID:3138286 A;Accession: A3513
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Molecule type: mRNA
A;Residues: 1-106 <AGA>
A;Residues: 1-106 <AGA>
A;Cross-references: UNIPARC:UPI00001767AC; GB:S68985; NID:g545746; PIDN:AAB30096.1;
A;Cross-references: UNIPARC:UPI00001767AC; GB:S68985; NID:g545746; PIDN:AAB30096.1;
A;Cross-references: UNIPARC:UPI0001767AC; GB:S68985; NID:g545746; PIDN:AAB30096.1;
A;Note: sequence inconsistent with nucleotide translation
A;Note: sequence extracted from NCBI backbone (NCBIN:144174, NCBIP:144175)
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: immunoglobulin homology <IMM>
                                                                        A;Cross-references: UNIPARC:UPI0000114D9C; GB:M20834; NID:g196943; C;Superfamily: immunoglobulin V region; immunoglobulin homology C;Keywords: heterotetramer; immunoglobulin P;38-113/Domain: immunoglobulin homology <IMM>
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F;71-77/Region: complementarity-determining 3
F;110-118/Region: complementarity-determining 3
F;130-140/Domain: complementarity-determining 3
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A; Residues: 1-130 < KOF>
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Best Local S
Matches 85
     Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         71.9%;
81.7%;
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78.4%;
  71.6%;
79.8%;

    Mismatches

 Mismatches

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Pred. No. 1e-29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 427.5;
Pred. No. 7.4
  Score 422.5; DB 2;
Pred. No. 1.8e-29;
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Ig light chain V region - mouse (fragment)
N,Alternate names: L6 anti-tumor antibody
C;Species: Mus musculus (house mouse)
C;Species: Mus musculus (house mouse)
C;Accession: A42848; 833902
R;Fell, H.P.; Gayle, M.A.; Yelton, D.; Lipsich, L.; Schieven, G.L.; Marken, J.S.
J. Biol. Chem. 267, 15552-1558, 1992
A;Title: Chimeric L6 anti-tumor antibody. Genomic construction, expression, and A;Reference number: A42848; MUID:92348410; PMID:1639794
A;Accession: A42848
A;Status: preliminary
A;Holecule type: DNA
A;Residues: 1-107 <FEL>
A;Cross-references: UNIPARC:UPI0000115334; EMBL:M90690
A;Cross-reference extracted from NCBI backbone (NCBIN:109958, NCBIP:109959)
C;Keywords: immunoglobulin region; immunoglobulin homology <IMM>
F;16-89/Domain: immunoglobulin homology <IMM>
                                                                                                                                                                                                                                                                                                                                                           J. Immunol. 142, 888-893, 1989
A;Title: Amino acid substitutions in V-H CDR2 change the idiotype
A;Reference number: A30562; MUID:89110066; PMID:2464031
                                                                                                                                                                                                                                                                                                                                                                                                   Ig kappa chain V regions (27.7.2 and 27.4b.2) - mouse C;Species: Mus musculus (house mouse) C;Date: 23-Mar-1989 #sequence_revision 23-Mar-1989 #te C;Accession: A30562 C;Accession: A30562 R;Sikder, S.K.; Borden, P.; Gruezo, F.; Akolkar, P.N.; Jimunol. 142, 888-893, 1989
                                                                                                                                                                                                 A/Cross-references: UNIPARC:UPI00001767B0 (5/Superfamily: immunoglobulin V region; immunoglobulin C/Keywords: heterotetramer; immunoglobulin P;16-89/Domain: immunoglobulin homology <11
                                                                                                                                                                                                                                                                                 A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-107 <SIK>
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75.7%;
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Pred. No. 5e-29;
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Pred. No. 2.8
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C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
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C; Date: 23-Mar-1989
C; Accession: B30562
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   R;Kofler, R.; Noonan, D.J.; Strohal, R.; Balderas, R.S.; Moller, N.P.H.; Dixon, Eur. J. Immunol. 17, 91-95, 1987
A;Title: Molecular analysis of the murine lupus-associated anti-self response: A;Reference number: S04573; MUID:87133856; PMID:3102255
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A; Residues: 1-107 < SIK>
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A; Residues: 1-130 < KOF>
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;Species: Mus musculus (house, mouse)
;Date: 23-Mar-1989 #sequence_revision
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                                                                                                            VPARFSGSGSGTSYSLTISRMEAEDAATYYCQQSNEDPYTFGSG 105
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Pred. No. 1.3e-28;
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Pred. No. 2.4e-28;
7; Mismatches 14;
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RESULT 11 S66536

Ig light chain V region - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 28-Oct-1996 #sequence\_revision 13-Mar-1997 #text\_change 20-Jun-2000

A;Title: Isolation and structural characterization of trimeric A;Reference number: S66536; MUID:95377318; PMID:7649183 A;Accession: S66536

cyanobacterial

photosystem

R;Tsiotis, G.; Haase, W.; Engel, A.; Michel, Eur. J. Biochem. 231, 823-830, 1995

C; Accession: S66536

A;Status: preliminary A;Molecule type: mRNA A;Residues: 1-120 <TS:

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C;Species: Mus musculus (house mouse)
C;Date: 06-Jan-1995 #sequence_revision 06-Jan-
C;Accession: S29590
R;Kavaler, J.
submitted to the EMBL Data Library, April 1991
A;Reference number: S28459
                                                                                                                                                                                                                                                                                                      R;Deng, J.B.; Han, H.; Su, C.Z.; Chen, C.O. Chinese Blochem. J. 12, 648-653, 1996
A;Title: Generation of a phage display library A;Reference number: PC4405
A;Contents: Spleen
                                                                                                                                                                                                                                                                                                                                                                                  pD0011
Ig kappa chain V region (VLB10, anti-AFP) - mouse (C;Species: Mus musculus (house mouse)
C;Date: 10-Jul-1998 #sequence_revision 10-Jul-1998
C;Accession: PD0011
C;Accession: PD0011
C;Accession: Pd0011
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A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-100 <KAV>
                                                                                                                                                                                                    A;Molecule type: mRNA
A;Residues: 1-107 < LDEN'>
A;Cross-references: UNIPARC:UPI00001767AA
C;Superfamily: immunoglobulin V region; immund
F;16-89/Domain: immunoglobulin homology < IMM>
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56 GVPAKGSGSGSTSYSLTISRVEAEDAATYYCQQWRDNPPTPGGGTKLEIKR
                              61 GVPARFSGSGSGTSYSLTISRMEAEDAATYYCQQSNEDPYTFGSGDQAGNKR
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                                                                                            1 DIELTQSPAIMSASPGEKVTTTCRASESVD9YGHSFMQWFQQXFGTSPKLWIYSTSNLAS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               IVLTQSPAIMSASPGEKVTMTCSASSSV-----SYMHWYQQKSGTSPKRWIYDTSKLASG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IELTQSPAIMSASPGEKVTTTCRASESVDSYGHSFMQWFQQKPGTSPKLWIYSTSNLASG 61
                                                                 Conservative
                                                                                                                                       Conservative
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77.9%;
                                                                                                                                                       69.1%;
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pred. No. 2.7e-28;
5; Mismatches 13;
                                                                                                                                       Score 407.5; DB 2;
Pred. No. 2.9e-28;
8; Mismatches 17;
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homology
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Ig kappa chain precursor V region (anti-phenyloxazolone 6F6) - mouse
C;Species: Mus musculus (house mouse)
C;Species: Mus musculus (house mouse)
C;Date: 31-Dec-1991 #sequence revision 09-Aug-1996 #text_change 16-Jul-1999
C;Accession: JL0079; A49044; MUID:95096973; PMID:31160
R;Kaartinen, M; Rocca-Serra, J; Maekelae, O.
Mol. Immunol. 25, 859-865, 1988
A;Fitle: Combinatorial association of V genes: one VH gene codes for three non-cross-real association of V genes: one VH gene codes for three non-cross-real A;Reference number: JL0079
A;Accession: JL0079
A;Accession: JL0079
A;Accession: JL0076; MUID:89096973; PMID:3211160
A;Accession: JL0079
A;Accession: JL0076; MUID:89096973; PMID:3211160
A;Accession: JL0079
A;Coss reference: UNIPARC:UPI000017371F; GB:M27792; NID:g197159
A;Coss reference: UNIPARC:UPI000017371F; GB:M27792;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Cross-references: UNIPARC:UPI000011760C; GB:S37663; NID:g250214; PIDN:AAB22331.1; A;Note: sequence extracted from NCBI backbone (NCBIN:106802, NCBIP:106809)
A;Notesion: B49044
A;Molecule type: DNA
A;Residues: 114-116 <MIL2>
A;Residues: 114-116 <MIL2>
A;Cross-references: UNIPARC;UPI000011E8B5; GB:S37664; NID:g250215; PIDN:AAB22332.1; A;Cross-references: BALB/C germ-line
A;Experimental source: BALB/C germ-line
A;Note: sequences extracted from NCBI backbone; partial nucleotide sequences of 13 a
A;Note: sequence extracted from NCBI backbone (NCBIN:106807, NCBIP:106822)
C;Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                    C/Superfamily: immunoglobulin V region, immunoglobulin homology C/Keywords: heterotetramer; immunoglobulin c/Keywords: heterotetramer; immunoglobulin predicted <SIG> F:122/Domain: signal sequence #status predicted <SIG> F:23-130/Product: Ig kappa chain V region (6F6) #status predicted F:38-111/Domain: immunoglobulin homology </MM>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Gene: V(kappa)Ox1
A;Introns: 17/1
C;Complex: An immunoglobulin heterotetramer subunit consists of two identical light (kap c;Complex: An immunoglobulin heterotetramer subunit consists of two identical light (kap hain disulfide bonds. In some cases, such as IgA and IgM, the subunits associate into la C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
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A; Residues: 1-25 <MIL>
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Matches 82; Conserv
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                                                                                                                                            IVLTQSPAIMSASPGEKVTMTCSASSSV-----SYMHWYQQKSGTSPKRWIYDTSKLASG
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       VPARFSGSGSGXSYSLTISSMEAEDAATYYCQQWSSNPLTFGAGTKLEXKR
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71.4%;
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Pred. No. 4.3e-28;
6; Mismatches 18
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Pred. No. 3.3e-28;
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                                    Ig kappa chain - mouse
C;Species: Mus musculus (house mouse)
C;Date: 19-Mar-1997 #sequence_revision 19-Mar-1997
C;Accession: S25058
R;Fischer, R.; Voss, A.; Niersbach, M.; Munziker, W
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S25058
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Ig kappa chain V region - mouse C,Species: Mus muscullus (house mouse) C,Date: 06-Jan-1995 #sequence_revision | C;Accession: S29591 R;Kavaler, J. submitted to the EMBL Data Library, Apr A;Reference number: $26459 A;Accession: $29591 A;Status: preliminary A;Molecule type: mRNA A;Residues: 1-103 < KNV>
                                                                                                                                                                                                                                              Ig kappa chain precursor V region - mouse
(;Species: Mus musculus (house mouse)
C;Becies: Ac-Oct-1989 #sequence_revision 26-Oct-1989 #text_change 21-Jan-2000
C;Accession: B32456
C;Accession: B32456
V. Biol. Chem. 264, 4513-4522, 1989
A;Title: Variable region primary structures of a high affinity anti-fluorescein immun A;Reference number: A32456; MUID:89174706; PMID:2494173
A;Accession: B32456
A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-130 <DOM>
A;Cross-references: UNIPARC:UPI00001148B9; GB:J04610; NID:g556313; PIDN:AAA50296.1; C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin homology <IMM*
F;38-113/Domain: immunoglobulin homology <IMM*
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S29591
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Best Local :
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Best Local
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83
                             64 ARPSGSGSGTSYSLTISRMEAEDAATYYCQQGNEDPYTFGSG 105
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                                                                                                                          4 LTQSPAIMSASPGEKVTTTCRASESVDSYGHBFMQWFQQKPGTSPKLWIYSTSNLASGVP
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Pred. No. 6.4e-28;
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Pred. No. 4.2e-28;
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EMBL Data Library,

M.; Munziker, July 1992

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A; Description: Production of a Tobacco mosaic virus (TMV) inactivating neotop specific # A; Reference number: $25057
A; Accession: $25058
A; Status: preliminary
A; Molecule type: mRNA
A; Residues: 1-235 <FIS>
A; Cross-references: UNIPARC: UPI0000116096; EMBL:X67211; NID:g54828; PIDN:CAA47650.1; PID C; Superfamily: immunoglobulin v region; immunoglobulin homology
F; 38-111/Domain: immunoglobulin homology <IMM>
Query Match
Best Local Similarity 74.8%; Pred. No. 1.2e-27;
Matches 83; Conservative 5; Mismatches 18; Indels 5; Gaps 1;
Oy 2 IELTQSPAIMSASPGEKVTTTCRASESVDSYGHSFMQWFQQKGTSPKLMIYSTSNLASG 61
Db 24 IVLTQSPAIMSASPGEKVTMTCSASSSV-----SKMQWFQQKSGTSPKRWIYDTSKLASG 78

Qy 62 VPARPSGSGSGTSYSLTISRMEAEDAATTYYCQQMSSUPLTFGAGTKLELKR 129
Db 79 VPGRPSGSGSGTSYSLTISRMEAEDAATTYYCQQMSSNPLTFGAGTKLELKR 129
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01-OCT-2002 (TrEMBLrel. 22, Created)
01-OCT-2002 (TrEMBLrel. 22, Last sequence update)
01-OCT-2002 (TrEMBLrel. 26, Last annotation update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Anti-VIPase light chain variable region (Fragment).
Mus musculus (Mouse).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia, Eutheria, Evarchontoglires; Glires; Rodentia; Sciurognat
                                                                                                                                                             Q8K1F3;
Q1-OCT-2002 (TrEMBLrel. 22, Created)
Q1-OCT-2002 (TrEMBLrel. 22, Last sequence update)
Q1-OCT-2004 (TrEMBLrel. 26, Last sequence update)
Q1-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Anti-VIPase light chain variable region (Fragment).
Mus musculus (Mouse).
Bukaryota, Metazoa, Chordata, Craniata, Vertebrata, Buteleostomi,
Mammalla, Butheria, Buarchontoglires, Glires, Rodentia, Sciurognat
Muridae, Murinae, Mus.
MCBI TaxID=10090;
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NUCLEOTIDE SEQUENCE.
STRAIN-BALB/c; TISSUE-Hyperimmunized spleen;
Zhou Y.-X., Taguchi H., Planque S., Karle S.,
Zhou Y.-X. (MAY-2002) to the EMBL/GenBank/DDBJ
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[3] 
NUCLECTIDE SEQUENCE.

MEDLINE=92381444; PubMed=1512540; DOI=10.1084/jem.176.3.761;

Tilman D.M., Jou N.T., Hill R.J., Marion TN.;

"Byth 1gM and 1gG anti-DNA antibodies are the products of clusterive B cell stimulation in (NZB x NZW)F1 mice.";

J. Exp. Med. 176:761-779(1992).
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PIR; A33933; A33933.
PIR; PH1058; PH1058.
HSSP; P01837; 25C8.
SMR; Q8K1F1; 2-114.
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Proc. Natl. Acad. Sci. U.S.A. 86:4624-4628(1989).
           NUCLEOTIDE SEQUENCE.
STRAIN-BALB/c; TISSUE-Hyperimmunized spleen;
Zhou Y.-X., Taguchi H., Planque S., Karle S.,
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InterPro; IPR003596; Ig-v.
SMART; SM00406; IGv; 1.
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RESULT 4
Q811C3_MOUSE PRELIMINARY;
AC Q811C3;
DT 01-JUN-2003 (TrEMBLrel. 24,
DT 01-JUN-2003 (TrEMBLrel. 24,
DT 01-JUN-2003 (TrEMBLrel. 26,
DT 01-MAR-2004 (TrEMBLrel. 26,
DT 01-MAR-2004 (TrEMBLrel. 26,
DE Imunoglobulin gamma-3 kappa
OS Mus musculus (Mouse).
CC Enkaryota, Metazoa; Chordate
OC Muridae; Murinae; Mus.
OC MINITERIALIS SEQUENCE.
RN NUCLEOTIDE SEQUENCE.
RR MUTHORIALIS SEQUENCE.
RR Takahashi S., Itoh J., Nosee
RT "Cloning and CDNA sequence:
RT Takahashi S., Itoh J., Nosee
RT "Cloning and CDNA sequence:
RT Takahashi S., Itoh J., Nosee
RT "Cloning and CDNA sequence:
RT HASP; P014C29; BAA03482.1;
DR HSSP; P014C29; BAA03482.1;
DR HSSP; P015C79; ZFBJ.
DR SMR; Q811C3; 23-131.
DR SMR; Q811C3; 23-131.
DR SMR; O81C3; 23-131.
DR SMR; O81C3; 23-131.
DR SMR; O81C3; 23-131.
DR SMR; S1GNAL 1 22
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                                                                                                                         XX MEDLINE-9315672; PubMed-842933; DOI=10.1016/0161-5890(93)90089-T;
XX MEDLINE-9315672; PubMed-842933; DOI=10.1016/0161-5890(93)90089-T;
XX Takahashi S., Itoh J., Nose M., Ono M., Yamamoto T., Kyogoku M.;
XY "Cloning and cDNA sequence analysis of nephritogenic monocional
XY antibodies derived from an MRL/lpr lupus mouse.";
XY Mol. Immunol. 30:177-182(1993).
XY EMBL; D14629; BAA03482.1; -; mRNA.
XY RSSP; P01679; 2FBJ.
XY HSSP; P01679; 2FBJ.
XY RSSP; P01679; 2FBJ.
XY RSSP; O811C3; 23-131.
XY SMR; O811C3; 23-131.
XY SMR; O811C3; 23-131.
XY SMR; O811C3; 23-131.
XY SMR; O811C3; 13-11.
XY SMR; O
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Best Local S
Matches 85
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"Both IgM and IgG anti-DNA antibodies are the products of selective B cell stimulation in (NZB x NZW)F1 mice.";

J. Exp. Med. 176:761-779 (1992).

EMBL, AF516282, AAM64200.1; -; Genomic_DNA.

PIR; A3393; A3393;

PIR; PH1085, PH1085.

HSSP, PO1837; 25C8.

SMR; Q8KIF9; 2-112.

Ensembl; ENSATTO-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Imunoglobulin gamma-3 kappa chain precurbor (Fragment). Mus musculus (Mouse). Bukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Buarchontoglires; Glires; Rodentia; Sciurognal Muridae; Mus.
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Interpro; IPR007110; Ig-like:
Interpro; IPR003596; Ig_v.
SMART; SM00406; IGv; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          nonmutated germ-line genes.";
Proc. Natl. Acad. Sci. U.S.A. 86:4624-4628(1989)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Baccala R., Quang T.V., Gilbert M., Ternynck T., Avr
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PubMed=2499887;
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3 (TrEMBLrel. 24,
4 (TrEMBLrel. 26,
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112 AA; 12017 MW;
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75.9%;
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Last annotation updat
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Pred. No. 4.6e-34;
6; Mismatches 16
                    Potential.
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                                                                                                                  Query Match
                                                                                                                                                  EMBL; AJ416331; CAC94866.1; -; mRNA.

PIR; G27887; G27887.

HSSP; P01834; 1MIM.

SMR; Q8VDD0; 23-134.

Ensembl; ENSMUSG0000062047; Mus musculus.

MGI; MGI:2686348; Gm1502.

InterPro; IPR007710; Ig-11ke.

InterPro; IPR007596; Ig_v.

SMART; SM00406; IGv; 1.

PROSITER; PS50835; IG_LIKE; 1.

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NON TER
SEQUENCE
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01-MAR-2002 (TrEMBLrel. 20, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Anti-MOG Z12 variable light chain (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Cranlata; Vertebrata; Buteleostomi;
Mammalia; Butheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;
Muridae; Murinae; Mus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mus musculus (Mouse).
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                                                                                                                                                                                                                                                                                                                                                                                                            STRAIN=BALB/c;
                                                                                                                                                                                                                                                                                                                                                                                                                                               Submitted (JAN-2002)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUCLEOTIDE SEQUENCE.
                                                                                                                                                                                                                                                                                                                        "Structural and functional implications
                                                                                                                                                                                                                                                                                                                                                PubMed=2427335;
                                                                                                                                                                                                                                                                                                                                                                                    Submitted (OCT-2001)
                                                                                                                                                                                                                                                                                                                                                                                                 Chernajovsky Y.;
                                                                                                                                                                                                                                                                                                                                                                                                                       NUCLEOTIDE SEQUENCE
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                                                                                                                                                                                                                                                                                               hemagglutinin.";
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                                                                                                        Similarity
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               VPARFSGSGSGTSYSLTISRMEAEDAATYYCQQSNEDPYTFGSGDQAGNKR 112
                                                                   IELTQSPAIMSASPGEKVTTTCRASESVDSYGHSFMQWFQQKPGTSPKLWIYSTSNLASG
                                             VPARPSGSGSGTSYSLTISSMEAEDAATYYCHQRSSYPWTFGGGTKLEIKR
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131 AA;
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Pred. No. 4.8e-33;
7; Mismatches 11
                                                                                           Score 411.5; DB 2; Length 134; Pred. No. 7e-33; 6; Mismatches 18; Indels 5
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on the influenza virus
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RX MEDLINE-22388257, PubMede12477932; DOI=10.1073/pnas.242603899;
RX MIDLINE-22388257, PubMede12477932; DOI=10.1073/pnas.242603899;
RX Alausner R.D., Collins F.S., Wagner L., Shanmen C.M., Schuler G.D.,
RA Altechul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Altechul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hong L.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Stapleton M.J., UEddin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Brownstein M.J., UEddin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Brownstein M.J., Wadin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Rodriguez A.C., Grimwood J., Schernan A., Rodrigues S., Sanchez A.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J.M., Marra M.A.;
RA Rodriguez A.C., Grimwood J., Schmutz J.M., Marra M.A.;
RA Rodriguez A.C., Grimwood J., Schmutz J.M., Marra M.A.;
RA Rodriguez A.C., Grimwood J., Schmutz J.M., M
                                                                                                                                                                                                                                                                                       Query Match
Best Local S
Matches 82
                                                                                                                                                                                                                                                                                                                                                                                                                                 InterPro; IPR003599; IG.

InterPro; IPR003597; IG_C1.

R InterPro; IPR003597; IG_C1.

R InterPro; IPR003597; IG_MIC.

R InterPro; IPR003596; IG_V.

R InterPro; IPR003596; IG_V.

R Pfam; PP07654; C1-set; 1.

R SWART; SM00409; IG; 2.

R SWART; SM00406; IGC1; 1.

R SWART; SM00406; IGV; 1.

R SWART; SM00406; IGV; 1.

R SWART; SM00406; IGV; 1.

R PROSITE; PS00290; IG_MIC; UNKNOWN_1.

D SEQUENCE 235 AA; 25719 MW; BE4E4ABDD2578252 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Q58EV6 MOUSE PRELIMINARY;
Q58EV6;
10-MAY-2005 (TrEMBLrel. 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mus musculus (Mouse).

Mus musculus (Mouse).

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;

Mammalla; Butheria; Buarchontoglires; Glires; Rodentia; Sciurognathi;

Muroidea; Muridae; Mus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Submitted (MAR-2005) to the EMBL/GenBank/DDBJ databases. EMBL; BC091738; AAH91738.1; -; mRNA. SMR; Q58EV6; 23-235.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUCLEOTIDE SEQUENCE.
STRAIN=FVB/N; TISSUE=Colon;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NCBI_TaxID=10090;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SMR; Q58EV6; 23-235.
GO; GO:0003823; F:antigen binding; IEA.
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   79
                                                                            62
                                                                                                                                                 24
                                                                                                                                                                                                                                                                                                                             Similarity
                                                                    VPARFSGSGSGTSYSLTISRMEAEDAATYYCOOSNEDFYTFGSGDQAGNKR 112
                                                                                                                                                                                           IELTQSPAIMSASPGEKVTTTCRASESVDSYGHSFMQWFQQKFGTSFKLWIYSTSNLASG
   VPARFSGSGSGTSYSLTISSMEAEDAATYYCQQWTSNPLTFGAGTKLDLKR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sci. U.S.A. 99:16899-16903(2002)
                                                                                                                                                                                                                                                                                                                                 68.7%;
73.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Last sequence update)
Last annotation update)
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                                                                                                                                                                                                                                                                                               5
                                                                                                                                                                                                                                                                                               Score 405.9; DB 2;
Pred. No. 9.4e-32;
5; Mismatches 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRT;
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RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Altschul S.F., Zeeberg B., Buctow K.H., Schaefer C.F., Bhat N.K.,
RA Altschul S.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Stapleton M., Oldin T.B., Bonaldo M.F., Casavant T.L., Scheetz T.S.,
RA Robanstein M.J., Uddin T.B., Bonaldo M.F., Carrinci P., Prange C.,
RA Rahà S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Rahà S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA Bosak S.A., McEwan P.J., McKernan K.J., Lu X., Gibbs R.A.,
RA Villalon D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,
RA Villalon D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,
RA Phely J., Helton E., Ketteman M., Madan A., Roorigues S., Sanchez A.,
RA Mhiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Schien J.E., Jones S.J.M., Marra M.A.,
The Control of the Schmutz J.M., Marra M.A.,
The Control of the Schmutz J.M. Marra M
                                                                                                                                                                                                                                                                                                        Matches
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Best Local
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Interpro; IPR003199; Ig-1ike.
Interpro; IPR003597; Ig_c1.
Interpro; IPR003597; Ig_c1.
Interpro; IPR003596; Ig_w:
Pfam; PP07564; C1-set; I.
SMART; SM00409; IG; 2.
SMART; SM00407; IGc1; 1.
SMART; SM00406; IGv; 1.
SMART; SM00406; IGv; 1.
PROSITE; P850835; IG_LIKE; 2.
PROSITE; P850290; IG_MEC; UNKNOWN 1.
SEQUENCE 237 AA; 25978 MW; A8859
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mus musculus (Mouse).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Buarchontoglires; Glires; Rodentia; Sciurognathi;
Muroidea; Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SMR, Q569Y8; 23-237.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Submitted (APR-2005) to the EMBL/GenBank/DDBJ databases. EMBL, BC092251; AAH92251.1; -; mRNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUCLECTIDE SEQUENCE.
STRAIN-FVB/N; TISSUE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRAIN=FVB/N; TISSUE=Kidney;
MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUCLEOTIDE SEQUENCE
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80
                                                   61 GVPARFSGSGSGTSYSLTISRMEAEDAATYYCQQSNEDPYTFGSGDQAGNKR 112
                                                                                                                                                  23
                                                                                                                                                                                      1 DIBLTQSPAIMSASPGEKVTTTCRASBSVDSYGHSFMQWFQQKPGTSPKLWIYSTSNLAS
                                                                                                                                                                                                                                                                                                                                            Similarity
GVPFRFSGSGSGTSYSLTISSMEAEDAATYYCQQMNYYPLTFGGGTRLELKR 131
                                                                                                                                          EIVLTQSPALMAASPGEKVTITCSVSSSIDS---SNLHWYQQKSGTSPKAWIYGTSNLAS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Acad. Sci. U.S.A. 99:16899-16903 (2002).
                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TISSUE=Kidney;
                                                                                                                                                                                                                                                                                                                                68.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Created)
Last sequence update)
Last annotation update)
                                                                                                                                                                                                                                                                                             Score 402.5; DB 2; Length 23 Pred. No. 1.1e-31; 7; Mismatches 21; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                           A88596AA47FFB932 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                            2; Length 237;
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RESULT 8

RESULT 9
KV3H\_MOUSE

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Q5XFY8 MOUSE PRELIMINARY;
AC G5XFY8 MOUSE PRELIMINARY;
AC G5XFY8;
DT Q55CT-2004 (TTEMBLIFE1 28,
DT 25-CCT-2004 (TTEMBLIFE1 28,
DT 25-CCT-2004 (TTEMBLIFE1 28,
DT 25-CCT-2004 (TTEMBLIFE1 28,
DT 25-CCT-2004 (TTEMBLIFE1 28,
DE Hypothetical protein.

OS Mus musculus (Mouse).

GE WARTYOLA, Metazoa; Chordato Communalia; Eutheria; Euarchoo
OC Muroidea, Muridae; Murinae;
RP MUCLEOTIDE SEQUENCE.
RP MUCLEOTIDE SEQUENCE.
RC STRAIN=FV9/N, TISSUE-Colon;
RX MEDLINE=2238857; PubMede-12.
RA Strausberg R.L., Feingold E RA Klausner R.D., Collins F.S.
RA Altachul S.F., Zeeberg B., S.
RA Altachul S.F., Zeeberg B., S.
RA Hopkins R.F., Jordan H., Mouralina K., Ra Hopkins R.S., Loquellano N.A., Ra Booak S.A., McEwan P.J., Md.
RA Booak S.A., McEwan P.J., Md.
RA Richards S., Worley K.C., H.
RA Villalon D.K., Madan A., Young RA Richards S., Worley K.C., H.
RA Villalon D.K., Madan A., Young RA Bakesley R.W., Touchman J.
RA Rodriguez A.C., Grimwood J.
RA ROGRIGUE SEQUENCE.
RC STRAIN=FWB/N; TISSUE-Colon;
RA GOLOGOSOS SEQUENCE
RC STRAIN=FWB/N; TISSUE-Colon;
RA Director McC Project;
RN []
RR MRHJ BC08463, AAR84683.1;
DR SMR, Q5XFWS; 33-235
DR GO; GO:0003823; F:antigen b
DR InterPro; IPR003599; Ig. C1.
DR SMART; SM00407; IGG-1; 1.
DR SMART; SM00406; IGy 1.
DR SMART; SM00406; IGy 1.
DR SMART; SM00406; IGM-C; U
KW Hypothetical protein.
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                                                                                                                                                                                                                                                                  Query Match
Best Local S
Matches 79
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRAIN=FVB/N; TISSUS=Colon;

Director MGC Project;

Blbmitted (OCT-2004) to the EMBL/GenBank/DDBJ databases.

RMBL; BC08463; AAH84683.1; -; mRNA.

RMBL; BC084633; Fantigen binding; IBA.

GO:0003823; Fantigen binding; IBA.

GITEPTO; IPR00359; IG.

InterPro; IPR00359; IG.

SMART; SM00406; IG.; 1.

SMART; SM00406; IG.; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;
Muroidea; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                              PROSITE; PS50835, IG LIKE; 2.
PROSITE; PS00290; IG_MHC; UNKNOWN_1.
Hypothetical protein.
SEQUENCE 235 AA; 25835 MW; 828E9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      25-OCT-2004 (TREMBLIRE1. 28, 25-OCT-2004 (TREMBLIRE1. 28, 25-OCT-2004 (TREMBLIRE1. 28,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
79
                                                            62
                                                                                                                               24
                                                                                                                                                                                                                                                                                                     Similarity
                                              VPARFSGSGSGTSYSLTISRMEAEDAATYYCQQSNEDPYTFGSGDQAGNKR 112
VPARFSGSGSGTSYSLTISRVEAEDAATYYCQQWSSNPLTFGAGTRLELKR
                                                                                                                               IVLSQSPAILSGFPGEKVIMICRASSSV-----NYMHWYQQKPGSSPKPWIYAISKLASG
                                                                                                                                                                                               IELTQSPAIMSASPGEKVTTTCRASESVDSYGHSFMQWFQQKPGT8PKLWIYSTSNLASG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TISSUE=Colon;
                                                                                                                                                                                                                                                                                                                                                                                                            25835 MW; 828E9ED277FDC667 CRC64;
                                                                                                                                                                                                                                                              68.1%; Score 401.5; DB 2; 71.2%; Pred. No. 1.3e-31; tive 12; Mismatches 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Created)
Last sequence update)
Last annotation update)
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RESULT 10
KV3J MOUSE
ID KV3J
AC P0166
DT 21-JU
DT 21-JU
DT 110-MA
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P01660;
21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence update)
10-MAY-2005 (Rel. 47, Last annotation update)
1g kappa chain V-III region PC 3741/TEPC 111.

Mus musculus (Mouse).

Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalla; Eutheria, Euarchontoglires, Glires, Rodentia, Sciurognat
Maroides, Muridae; Muridae; Mus.
MOUSE P01662;
P01662;
21-JUL-1986
21-JUL-1986
11-MAY-2005
                                                                                                                                                                                                                                                                                                                                                       IG LIKE;
Immunoglobulin V region
REGION
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REGION
REGION
REGION
REGION
DISULFID
NON TER
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDIJINE-79012520; PubMed-99744;
MCKean D.J., Bell M., Potter M.;
McKean B.J., Bell M., Potter M.;
Mechanisms of antibody diversity: multiple genes encode s
related mouse kappa variable regions.";
Proc. Natl. Acad. Sci. U.S.A. 75:3913-3917(1978).
-!- MISCELLANEOUS: The PC 3741 and TEPC 111 sequences are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PROTEIN SEQUENCE (PC 3741).
MEDLINS=79073152; PubMed=103003;
Weigert M., Gatmaitan L., Loh B.,
"Rearrangement of genetic informa"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This Swiss-Prot entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the El the European Bioinformatics Institute. There are no rest
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nature 276:785-790(1978).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             diversity.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PROTEIN SEQUENCE (TEPC 111).
MEDLINE=79012520; PubMed=99744;
                                                                                                                                                                                                                                                                                                                                                                                                                                              Ensembl; ENSMUSG00000060064;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PIR; A93204; KVMS37.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             removed
                                                                                                                                                                                                                                                                                                                                                REGION
                                                                                                                                                                                                                                                                                                                                                                                                                    InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            HSSP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         as long
                                                                                                                                                                                                                                                                                                                                                                                                                                                            P01665;
                                                                                                                                 61
                                                                                                           13
                                                                                                                                                                                                                                                                                                                                                                                               SM00406; IGV; 1.
E; PS50835; IG_LIKE;
                                                                                                                                                         -
                                                                                                                                                                              \vdash
                                                                                                                                                                                                                   Similarity
                                                                                                                                                                     DIELTQSPAIMSASPGEKVTTTCRASESVDSYGHSFMQWFQQKPGTSPKLWIYSTSNLAS
                                                                                                                           GVPARFSGSGSGTSYSLTISRMEAEDAATYYCQQSNEDPYTFGSG 105
                                                                                                           GIPARPSGSGSRTDFTLTINPVEADDVATYYCOQSNEDPYTFGGG
                                                                                                                                                         DIVLTOS PASLAVSLGQRATISCRASES VDS YGNS PMHWYQQKPGQPPKLLIYRASNLES
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102
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  (Rel.
(Rel.
(Rel.
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                                                 STANDARD;
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01, Last sequence update)
47, Last annotation updat
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information
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Immunoglobulin domain;
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                                                                                                                                                                                                                  Score 401; DB 1;
Pred. No. 6.2e-32;
                                                                                                                                                                                                                                                                                        Framework-3.
Complementarity-determining-3
Framework-4.
                                                                                                                                                                                                                                                                                                                        Complementarity-determining-1. Framework-2. Complementarity-determining-2.
                                                                                                                                                                                                                                                                                                                                                               Framework-1
                                                 PRT;
                                                                                                                                                                                                                                                     EC46C9D259213BE4
                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                            similarity.
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                                                                                                                                                                                                                                                                                                                                                                                                                                               musculus
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tion may produce immunoglobulin
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                                                 111
                                                 B
                                                                                                                                                                                                                             Length 111,
                                                                                                                                                                                                                                                      CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      identical.
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RESULT
Q9U410
ID Q9
AC Q9
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Matches 74
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PROTEIN SEQUENCE (PC 9245).

MEDLINE=79073152; PubMed=103003;

MEDLINE=79073152; PubMed=10 10 B., Schilling

Annetic information may
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Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;
Muroidea; Muridae; Musimae; Mus.
                                                                                                                                                                                                                                                                                                                                   REGION
DISULFID
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SEQUENCE
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MEDLINE=79012520; PubMed=99744;
MCKean D.J. Bell M., Potter M.;
"Mechanisms of antibody diversity: multiple
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ig kappa chain V-III region ABPC 22/PC Mus musculus (Mouse)
                                    01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
Monoclonal anti-idiotypic Schistosoma japonicum antibody NP30
immunoglobulin light chain variable region (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
SMART; SM00406; IGv; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This Swiss-Prot entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation - the European Bioinformatics Institute. There are no restrictions on its
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               related mouse kappa variable regions "; FPPP proc. Natl. Acad. Sci. U.S.A. 75:3913-3917(1978).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NCBI_TaxID=10090;
            Mus musculus (Mouse).
Eukaryota; Metazoa; C
                                                                                                      Q9U410;
                                                                                                                Q9U410_MOUSE PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PROSITE; PS50835; IG LIKE;
Direct protein sequencing;
Immunoglobulin V region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ensembl; ENSMUSG00000060064; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SMR; P01662;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nature 276:785-790(1978).
-!- MISCELLANEOUS: The ABPC22 and PC9241 sequences are identical.
Mammalia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     REGION
                                                                                                                              MOUSE
                                                                                                                                                                                 2
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                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                              NIVLTOS PASLAVSLGQRATIS CRASES VD9 YGNS FMHWY QQKPGQPPKLLIYLASNLES
                                                                                                                                                                                                                                                DIELTQSPAINSASPGEKVTTTCRASESVD$YGHSFMQWFQQKPGTSPKLWIYSTSNLAS
Butheria; Euarchontoglires;
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  Chordata; Craniata; Vertebrata; Euteleostomi; Euarchontoglires; Glires; Rodentia; Sciurogna
                                                                                                                                                                                                                                                                                                                                       12041 MW;
                                                                                                                                                                                                                                                                                                67.1%;
                                                                                                                                                                                                                                                                                    13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Immunoglobulin
                                                                                                                                                                                                                                                                                                Score 396; DB 1;
Pred. No. 1.9e-31;
                                                                                                                                                                                                                                                                                                                                                                                                     Complementarity-determining-2. Framework-3.
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                                                                                                                                                                                                                                                                                                                                                               By similarity.
                                                                                                                                                                                                                                                                                                                                      D7DF0609303453CE CRC64;
                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 musculus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Schilling J., Hood L.E.;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   domain;
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    Sciurognathi;
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Best Local S
Matches 79
                                                                                                                                                                                                                                                                                       P04940;
13-AUG-1987 (Rel. 05, Created)
13-AUG-1987 (Rel. 05, Last sequence update)
10-MAY-2005 (Rel. 47, Last annotation update)
10-MAY-2005 (Rel. 47, Last annotation update)
19 kappa chain V-VI region NQ2-17.4.1.
Mus musculus (Mouse).

Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Bukaryota; Mutheria; Euarchontoglires; Glires; Rodentia; Sciurognat
Murridae; Muridae; Mus.
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Song X.T., Feng Z.Q., Qiu Z.N., Li Y.Q., Yu X.C., Xiong Y., Yin C., Song H.L., Guan X.H.;
Huang H.L., Guan X.H.;
"[Cloning and sequence analysis of the light chain variable region gene of monoclonal anti-idiotypic antibody NP30 of Schistosoma japonicum].";
                             SMR; P04940; 1-107.
Ensembl; ENSMUSG0000062047; Mus musculus.
Interpro; IPR007710; Ig-11ke.
Interpro; IPR003596; Ig_v.
                                                                                                                                          This Swiss-Prot entry is copyright. It is produced through a obstween the Swiss Institute of Bioinformatics and the EMBL the European Bioinformatics Institute. There are no restrict
                                                                                                                                                                                                                                                                                                                                                                                                                       MOUSE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SMR; Q9U410; 4-106.
InterPro; IPR007110; Ig-11)
InterPro; IPR003596; Ig_v.
SMART; SM00406; IGv; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Zhongguo Ji Sheng Chong Xue Yu
18:257-259(2000).
EMBL; AF207620; AAF19434.1; -;
                         SMART; SM00406; IGv;
                                                                                                                                                                                                                                                                                     NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                        KV6F_MOUSE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Muridae; Murinae; Mus
               PROSITE; PS50835;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PROSITE; PS50835; IG_LIKE; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        HSSP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUCLEOTIDE SEQUENCE
                                                                                                                                                                                          - FUNCTION: Anti-2-phenyl oxazolone (PHOX) Antibody.
                                                                                   K00735; AAA38680.1; -; P01679; 2FBJ.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  59 VRFSGSGSGTSYSLTISRMEAEDAATYYCQQWTSYPFTFGSG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      P01679; 2FBJ.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                64 ARFSGSGSGTSYSLTISRMEAEDAATYYCQQSNEDPYTFGSG 105
                                                                                                                                long as its
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4 LTQSPAIMSASPGEKVTTTCRASESVDSYGHSFMQWFQQKPGTSPKLWIYSTSNLASGVP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LTQSPAIMSASPGEKVTMTCSASSSV-----SYVYWYLQKPGSSPRLLIYDTSNLASGVP
1mmunoglobulin domain;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            106 AA;
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77.5%;
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                                                                                                mRNA
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Pred. No. 2.1
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  Immunoglobulin V region.
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                                                                                                                                  modified
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                                                                                                                                  and this statement
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                                                                                                                                              restrictions
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     106;
                                                                                                                                                                   a collaboration
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                                                                                                                                                                                                                               6
                                                                                                                                                        outstation
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RESULT 13

KV6G MOUSE
ID KV6G MA
AC P04941
DT 13-AUG
CC MARNIA
RT MARNIA
DR HSSP;
DR SMART;
DR EMBL;
DR EMBL;
DR TINTERF
DR TINTERF
DR TREGION
FT REGION
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13-AUG-1987 (Rel. 05, Created)
13-AUG-1987 (Rel. 05, Last sequence update)
10-MAY-2005 (Rel. 47, Last annotation updat
-- ranna chain V-VI region NQ2-48.2.2.
                  REGION
DISULFID
NON_TER
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Kaarrinen M., Griffiths G.M., Markham A.F., Milstein C.;
"mRNA sequences define an unusually restricted IgG respoi
phenyloxazolone and its early diversification.";
Nature 304,320-324(1983)
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InterPro; IPR003596; Ig_v.
SMART; SM00406; IGv; 1.
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-I- FUNCTION: Anti-2-phenyl oxazolone (PHOX) Antibody.
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21-JUL-1986 (Rel. 01, Last sequence up
10-MAY-2005 (Rel. 47, Last annotation
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MCKean D.J., Bell M., Potter M.;
McKean D.J., Bell M., Potter M.;
Mechanisms of antibody diversity: multiple genes encode
"Mechanisms of antibody diversity: multiple genes encode
related mouse kappa variable regions.";
Proc. Natl. Acad. Sci. U.S.A. 75:3913-3917 (1978).
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InterPro; IPR003596; Ig_v.
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                                                                                                        Score 391; DB 1; Pred. No. 7.4e-31;
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Pramework-1
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2: /cgn2_6/ptodata/1/iaa/H_COMB.pep:*
3: /cgn2_6/ptodata/1/iaa/H_COMB.pep:*
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5: /cgn2_6/ptodata/1/iaa/RE_COMB.pep:*
6: /cgn2_6/ptodata/1/iaa/AB_COMB.pep:*
    GenCore version 5.1.6 Copyright (c) 1993 - 2005 Compugen Ltd
    DB
US-09-798-689-8
US-09-798-689-21
US-09-171-945-9
US-09-171-945-17
US-09-10-059-17
US-09-10-059-17
US-09-10-059-17
US-09-138-2
US-08-116-7788-2
US-08-438-562-2
US-08-438-562-2
US-08-438-522-240
US-09-136-522-240
US-09-136-157-12
US-09-136-157-12
US-09-136-157-2
US-09-136-16-22-244
US-09-196-522-244
US-09-196-522-244
US-09-196-16-16
US-09-176-116-16
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 Sequence 8, Appli
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Sequence 240, Appl
Sequence 12, Appl
Sequence 13, Appl
Sequence 16, Appl
Sequence 237, App
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Query Best Match	APPLICANT: TITLE OF J	45 429 RESULT 1 US-09-798-68 ; Sequence 8 ; Patent No.	,	33000
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77.4%; Score 456.5; DB 2; Length 106; imilarity 85.7%; Pred. No. 1.6e-37; ; Conservative 3; Mismatches 7; Indels 5; Gaps DIELTQSPAINSASPGEKVITTCRASESVD9YGHSFMQWFQQKPGTSPKLWIYSTSNLAS		8 1		
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re 456.5; d. No. 1.6 Mismatches M	Reducing th Radia gs 1-41 9/798,68 1,163 7,113 6,804 6,533 6,552 6,041	08-477-531 ALIGNMENTS 9	4777 4777 4774 4887 4887 4777 4777	-423 -423 -469 -634
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RESULT 2 US-09-798-689-21 ; Sequence 21, Application US/09798689 ; Patent No. 6811779

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GENERAL INFORMATION:

APPLICANT: Emery, Stephen

APPLICANT: Copley, Clive Graham

APPLICANT: Copley, Clive Graham

APPLICANT: Edge, Michael Derek

TITLE OF INVENTION: Monoclonal Antibody to CEA, Conjugates Comprising Said

TITLE OF INVENTION: Antibody, and Their Therapeutic Use in an Adept System

FILE REFERENCE: Monoclonal Antibody to CEA

CURRENT APPLICATION NUMBER: U9/09/171,945

CURRENT APPLICATION NUMBER: U9/09/171,945

CURRENT FILING DATE: 1998-10-29

PRIOR APPLICATION NUMBER: GB9703103.3

PRIOR FILING DATE: 1997-02-14

PRIOR APPLICATION NUMBER: GB9609405.7

PRIOR APPLICATION NUMBER: PCT/GB97/01165

PRIOR APPLICATION NUMBER: PCT/GB97/01165

PRIOR APPLICATION NUMBER: PCT/GB97/01165

PRIOR FILING DATE: 1997-04-29

NUMBER OF SEQ ID NOS: 131

GORTMADER: SECONDATE: 131
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CURRENT FILING DATE: 2001-03-02
PRIOR APPLICATION NUMBER: 09/401,163
PRIOR PILING DATE: 1999-09-22
PRIOR APPLICATION NUMBER: 08/967,113
PRIOR PILING DATE: 1997-11-10
PRIOR PILING DATE: 1996-09-03
PRIOR PILING DATE: 1996-09-03
PRIOR PILING DATE: 1996-09-03
PRIOR APPLICATION NUMBER: 08/476,533
PRIOR APPLICATION NUMBER: 08/476,533
PRIOR PILING DATE: 1995-06-07
PRIOR PILING DATE: 1995-06-07
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION
                        Query Match
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ORGANISM: Mouse
-09-798-689-21
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PRIOR APPLICATION NUMBER: 08/326,552
PRIOR FILING DATE: 1994-10-20
PRIOR APPLICATION NUMBER: 08/196,041
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                                                                                                                                            LENGTH: 108
TYPE: PRT
ORGANISM: Artificial Sequence
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85.7%;
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Pred. No. 8e-37;
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APPLICANT: Copley, Clive G
APPLICANT: Edge, Michael Derek
APPLICANT: Edge, Michael Derek
APPLICANT: Emery, Stephen Charles
TITLE OF INVENTION: Monoclonal Antibody to CEA, Conjugates Comprising Said Antibody, &
TITLE OF INVENTION: Their Therapeutic use in an Adept System
FILE REFERENCE: 1991-209
CURRENT APPLICATION NUMBER: US/09/910,059
CURRENT APPLICATION NUMBER: US/09/910,059
CURRENT FILING DATE: 2001-07-23
PRIOR APPLICATION NUMBER: US/09/171,945
PRIOR FILING DATE: 1998-10-29
PRIOR APPLICATION NUMBER: PCT/GB97/01165
PRIOR APPLICATION NUMBER: PCT/GB97/01165
PRIOR APPLICATION NUMBER: BS 9703103.3
PRIOR APPLICATION NUMBER: GB 9609405.7
PRIOR APPLICATION NUMBER: GB 9609405.7
PRIOR FILING DATE: 1997-02-14
NUMBER OF SEQ ID NOS: 131
SOFTMARE: PatentIn version 3.1
APPLICANT: Emery, Stephen
APPLICANT: Copley, Cilve Graham
APPLICANT: Edge, Michael Derek
ITILE OF INVENTION: Monoclonal Antibody to CEA, Conjugates Comprising Said
TITLE OF INVENTION: Monoclonal Antibody to CEA
TITLE OF INVENTION: Antibody, and Their Therapeutic Use in an Adept System
FILE REFERENCE: Monoclonal Antibody to CEA
CURRENT APPLICATION UNMBER: US/09/171,945
CURRENT FILING DATE: 1998-10-29
PRIOR APPLICATION NUMBER: GB9703103.3
PRIOR FILING DATE: 1997-02-14
PRIOR APPLICATION NUMBER: GB9609405.7
PRIOR APPLICATION NUMBER: GB9609405.7
PRIOR PILING DATE: 1995-05-04
PRIOR FILING DATE: 1995-05-04
PRIOR FILING DATE: 1997-04-29
NUMBER OF SEQ ID NOS: 131
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Patent No. 6277599
GENERAL INFORMATION:
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Patent No. 6903203
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Local Similarity 80.4%;
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CURRENT PILLING DATE: 2001-07-23
PRIOR APPLICATION NUMBER: US 09/171,945
PRIOR PILING DATE: 1998-10-29
PRIOR APPLICATION NUMBER: PCT/GB97/01165
PRIOR PILING DATE: 1997-04-29
PRIOR APPLICATION NUMBER: GB 9703103.3
PRIOR PILING DATE: 1997-02-14
PRIOR APPLICATION NUMBER: GB 9609405.7
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Patent No. 6903203
GENERAL INFORMATION:
APPLICANT: Copley, Clive G
APPLICANT: Edge, Michael Derek
APPLICANT: Emery, Stephen Charles
TITLE OF INVENTION: Monoclonal Antibody to CEA, Conjugates Comprising Said Antibody,
TITLE OF INVENTION: Their Therapeutic use in an Adept System
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQ ID NO 17
LENGTH: 235
                                      GENERAL INFORMATION:
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                                                         Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION: chimaeric light chain sequence
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SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR FILING DATE: 1996-05-04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILE REFERENCE: 1991-209
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RAMOT UNIVERSITY AUTHORITY FOR APPLIED RESEARCH & INDUSTRIAL DVELOPMENT LTD.
                                                                                                                                                                                              GVPARFSGSGSGTSYSLTISRMEAEDAATYYCQQSNEDPYTFGSGDQAGNKR 112
                                                                                                                                                                                                                                                      DIELTQSPAIMSASPGEKVTITCSASSSV-----TYMHWFQQXPGTSPKLWIYSTSNLAS
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                                                                         Application US/09526738A
                                                                                                                                                                                                                                                                                                                                 Conservative
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Pred. No. 2e-36;
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RESULT 9 US-08-116-778E-2

Sequence 2, Application US/08116778E Patent No. 5830470

GENERAL INFORMATION:

APPLICANT:
APPLICANT:

NAKAMURA, KAZUYASU KOIKE, MASAMICHI SHITARA, KENYA HANAI, NOBUO

APPLICANT: KUWANA, YOSHIHISA APPLICANT: HASEGAWA, MAMORU TITLE OF INVENTION: HUMANIZED ANTIBODIES

49

NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:

3: NIXON & VANDERHYB P.C. 1100 NORTH GLEBB ROAD

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; TYPE: PRT
; ORGANISM: Humanus
US-09-526-738A-4
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LENGTH: 25
TYPE: PRT
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SOFTWARE: PatentIn version 3.1
SEQ ID NO 4
LENGTH: 258
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                                                                                                                                                                                                                                                                                                                                          FILE REFERENCE: 1196336
CURRENT APPLICATION NUMBER: US/09/526,738A
CURRENT FILING DATE: 2000-03-16
                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: RAMOT UNIVERSITY AUTHORITY FOR APPLIED RESEARCH & INDUSTRIAL DVELOPMENT APPLICANT: LTD.
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SOFTWARE: PatentIn version 3.1
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                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: SINGLE CHAIN ANTIBODY AGAINST MUTANT P53
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189
                                                                     134 DIELTOSPAIMSASPGEKVTITCSASSSV----NYMHWFQQKPGTSPKLWISSTSNLAS
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                  61 GVPARFSGSGSGTSYSLTISRMEAEDAATYÝCQQSNEDPYTFGSGDQAGNKR 112
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                                                                                                                                           Similarity 80.4 90; Conservative
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GVPARFSGSGSGTSYSLTISRMEAEDAATYYCOORSSYPYTFGGGTKLQIKR
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                                                                                                                                                             76.0%;
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Pred. No. 2.8e-36;
4; Mismatches 13
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                                                                                                                                                             Score 448.5; DB 2;
Pred. No. 2.8e-36;
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COUNTRY: U.S.A.

22201-4714

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RESULT 10
US-08-438-562-2
! Sequence 2, Application US/08438562
! Patent No. 5874255
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Matches
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INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 129 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/08/116
PILING DATE: 07-SEP-93
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: WILSON, MARY J.
REGISTRATION NUMBER: 32,955
REFERENCE/DOCKET NUMBER: 249-
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                  NAME/KEY: domain
LOCATION: 88..96
IDENTIFICATION METHOD:
IDENTIFICATION METHOD:
IDENTIFICATION METHOD:
OTHER INFORMATION: /pro
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IDENTIFICATION METHOD:
IDENTIFICATION METHOD:
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OTHER INFORMATION: /pro
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LOCATION: -22...1
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                                                                                                                                                    62 VPARFSGSGSGTSYSLTISRMEAEDAATYYCQQSNEDPYTFGSGDQAGNKR 112
                                                                                                                                                                                                           24 IVLTQSPAIMSASPGEKVTITCSASSSV----SYMHWFQQKPGTSPKLWIYSTSNLASG
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                                                                                                                                                                                                                                                                                         90; Conservative
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                                                                                                                                VPARFSGSGSGTSYSLTISRMEAEDAATYYCQQRSSYPYTFGGGTKLEIKR 129
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81.1%;
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D: WITH KNOWN SEQUENCE OR TO AN ESTABLISHED DI: CONSENSUS
/product= "HYPERVARIABLE REGION 3"
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CONSENSUS
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Pred. No. 1.5e-36;
3; Mismatches 13;
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US-08-438-562-2
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INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 129 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/438,562
PILING DATE: 10-MAY-95
CLASSIFICATION DATA:
APPLICATION DATA:
APPLICATION UMBER: 08/116,778
PILING DATE: 07-SEP-93
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: WILSON, MARY J.
PROTEDER TOWN MINHERY J. 32 955
Query Match 75.8%;
Best Local Similarity 81.1%;
Matches 90; Conservative
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APPLICANT:
APPLICANT:
APPLICANT:
                                                                                            NAME/KEY: domain
LOCATION: 88.96
IDENTIFICATION METHOD:
IDENTIFICATION METHOD:
IDENTIFICATION METHOD:
OTHER INFORMATION: /pro
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REFERENCE/DOCKET NUMBER: 24
TELECOMMUNICATION INFORMATION:
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CORRESPONDENCE ADDRESS:
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TITLE OF INVENTION: HUMANIZED ANTIBODIES
                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: domain
LOCATION: 24..33
IDENTIFICATION METHOD:
IDENTIFICATION METHOD:
IDENTIFICATION METHOD:
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LOCATION: -22..-1
IDENTIFICATION METHOD:
IDENTIFICATION METHOD:
IDENTIFICATION METHOD:
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                                                                                                                                                                                                                                                                                                                   NAME/KEY:
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(703)816-4100
Th NO: 2:
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KOIKE, MASAMICHI
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49..55
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DD: WITH KNOWN SEQUENCE OR TO AN ESTABLISHED
DD: CONSENSUS
/product= "HYPERVARIABLE REGION 3"
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D: WITH KNOWN SEQUENCE OR TO AN ESTABLISHED
D: CONSENSUS
/product= "HYPERVARIABLE REGION 1"
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CONSENSUS "HYPERVARIABLE REGION 2"
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Score 447.5; DB 1;
Pred. No. 1.5e-36;
3; Mismatches 13;
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GENERAL INFORMATION:
APPLICANT: NAKAMURI
APPLICANT: KOIKE,
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NUMBER OF SEQUENCES: 103
CORRESPONDENCE ADDRESS:
ADDRESSE: NIXON & VANDERHYE P.C.
STREET: 1100 NORTH GLEBE ROAD
CITY: ARLINGTON
STATE: VIRGINATION
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INFORMATION FOR SEQ ID NO: 92:
SEQUENCE CHARACTERISTICS:
LENGTH: 129 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ZIP: 22201-4714
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOCTWARE PATENTIN Release #1.0, Version
CURRENT APPLICATION DATA:
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LOCATION: -22...1
IDENTIFICATION METHOD:
IDENTIFICATION METHOD: B:
IDENTIFICATION METHOD: E:
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LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 129 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                      NAME/KEY:
LOCATION:
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IDENTIFICATION METHOD:
IDENTIFICATION METHOD:
IDENTIFICATION METHOD:
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IDENTIFICATION METHOD:
IDENTIFICATION METHOD:
IDENTIFICATION METHOD:
                                                          NAMB/KBY:
LOCATION: 88..96
IDENTIFICATION METHOD:
IDENTIFICATION METHOD:
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; IDENTIFICATION METHOD:
; OTHER INFORMATION: /pr
US-08-483-528B-92
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APPLICANT: Jackson, Ronald
APPLICANT: Holiger, Kasper
APPLICANT: Holiger, Kasper
APPLICANT: Clackson, Timothy
APPLICANT: Clackson, Timothy
APPLICANT: Clackson, Timothy
APPLICANT: Winter, Gregory
APPLICANT: Winter, Gregory
APPLICANT: Bonert, Timothy
TITLE OF INVENTION. Methods for Producing Members of
FILE REFERENCE: 213839-00013
CURRENT APPLICATION NUMBER: US/09/726,219A
CURRENT APPLICATION NUMBER: GB 9015198.6
PRIOR APPLICATION NUMBER: GB 9015198.6
PRIOR APPLICATION NUMBER: GB 9029845 3
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US-09-726-219A-240
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                                                                                   FEATURE:

OTHER INFORMATION: VL of scFv from mouse immunized with 2-phenyl-5-oxazolone US-09-726-219A-240
                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR FILING DATE: 1990-10-19
PRIOR PELICATION NUMBER: GB 9022845.3
PRIOR FILING DATE: 1990-10-19
PRIOR APPLICATION NUMBER: GB 9024503.6
PRIOR PILING DATE: 1990-11-12
PRIOR PELICATION NUMBER: GB 9104744.9
PRIOR PILING DATE: 1991-03-06
PRIOR APPLICATION NUMBER: GB 9110549.4
PRIOR APPLICATION NUMBER: GB 9110549.4
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                                                                                                                                                                                           NUMBER OF SEQ ID NOS: 272
SOFTWARE: Patentin version 3.1
SEQ ID NO 240
LENGTH: 108
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                    Query Match
Best Local Similarity
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APPLICANT: Cambridge Antibody Technology Limited
APPLICANT: Medical Research Council
APPLICANT: McCafferty, John
APPLICANT: Pope, Anthony
                                                                                                                                                                                                                                                                                                  PRIOR FILING DATE: 1991-05-15
PRIOR APPLICATION NUMBER: PCT/GB91/01134
PRIOR APPLICATION STREET 1991-07-10
PRIOR PPLICATION NUMBER: US 07/971,857
PRIOR PILING DATE: 1993-01-08
PRIOR APPLICATION NUMBER: US 08/484,893
                                                                                                                                                                                                                                                                              PRIOR FILING DATE: 1995-06-07
                                                                                                                                  LENGTH: 108
TYPE: PRT
ORGANISM: Artificial Sequence
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Griffiths, Andrew
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81.1%;
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Pred. No. 1.5e-36;
Score 442.5; DB 2;
Pred. No. 3.9e-36;
4; Mismatches 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
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                                          Length 108;
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         Indels
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Matches

Conservative

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Gaps

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55 60

DIELTOSPAIMSASPGEKVTTTCRASESVDSYGHSFMQWFQQKFGTSPKLWIYSTSNLAS

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APPLICANT: Winter, Gregory
APPLICANT: Bonert, Timothy
ITITE OF INVENTION: Methods for Producing Members of Specific Binding Pairs
FILE REFERENCE: 213839-00004
CURRENT FILING DATE: 1998-11-28
CURRENT FILING DATE: 1998-11-28
PRIOR APPLICATION NUMBER: GB 9015198.6
PRIOR FILING DATE: 1990-07-10
PRIOR FILING DATE: 1990-07-10
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR APPLICATION NUMBER: GB 9024503.6
PRIOR FILING DATE: 1990-10-19
PRIOR APPLICATION NUMBER: GB 9024503.6
PRIOR APPLICATION NUMBER: GB 9024503.6
PRIOR APPLICATION NUMBER: GB 9024503.6
PRIOR APPLICATION NUMBER: GB 9104744.9
PRIOR APPLICATION NUMBER: GB 9104744.9
PRIOR PILING DATE: 1990-03-06
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                                                                                                                                                                                                                                                     ; OTHER INFORMATION: VL of scFv from mouse immunized with 2-phenyl-5-oxazolone US-09-196-522-240
                                                                                                                                                                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: Pa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Cambridg
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APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
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atent No. 6916605
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                                                                                                                                                                                                                                                                                             TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                LENGTH: 108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 1993-01-08
APPLICATION NUMBER: US 08/484,893
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: PCT/GB91/01134
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: GB 9110549.4 FILING DATE: 1991-05-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US 07/971,857
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            56 GVPTRFSGSGSGTSYSLTISRMEAEDAATYYCQQRSSYPPTFGSGTKLBIKR 107
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                                                                                                                         1 DIELTQSPAIMSASPGEKVTTTCRASESVDSYGHSFMQWFQQKPGTSPKLWIYSTSNLAS
                                                                                                                                                                                                                                                                                                                                                                                                     PatentIn version 3.1
                          GVPARFSGSGSGTSYSLTISRMEAEDAATYYCQQSNEDPYTFGSGDQAGNKR 112
GVPTRFSGSGSGTSYSLTISRMEAEDAATYYCQQRSSYPPTFGSGTKLEIKR 107
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Griffiths, Andrew
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            McCafferty, John
                                                                                                                                                                   Conservative
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79.5%;
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Pred. No. 3.9e-36;
                                                                                                                                                                 Mismatches
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                                                                               -NYMHWFQQKPGTSPKLWIYRTSNLAS
                                                                                                                                                                                                         Length 108;
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US-08-434-000A-12
RESULT 15
US-09-312-157-12
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
NAME: Guise, Jeffrey W.
REGISTRATION NUMBER: 34,613
REFERENCE/DOCKET NUMBER: 212/
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 552-8400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08
PILING DATE: 12/30/94
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/
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MEDIUM TYPE: 3.5" Diskette,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: ANDREW C. HIATT, JULIAN
APPLICANT: K.-C. MA, THOMAS LEHNER
TITLE OF INVENTION: IMMUNOGLOBULING CONTAINING PROTECTION
TITLE OF INVENTION: PROTEINS IN PLANTS AND THEIR USES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                         TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                        LENGTH: 105 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                           TELEX: 67-3510
TELEX: SEQUENCE LISTING
                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 90071
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STATE: California
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OPERATING SYSTEM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMATION:
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                                                                                    61 GVPARFSGSGSGTSYSLTISRMEAEDAATYYČQQSNEDPYTFGSG 105
                                                                                                                                                                                                      ,
96
                                                                                                                                                                                                                        Similarity
                                                                  GVPARFSGSGSGTSYSLTISRMBABDAATYYCHORTSYPYTFGGG
                                                                                                                                                          DIELTQSPAIMSASPGEKVTTTCRASESVDSYGHSFMQWFQQKPGTSPKLWIYSTSNLAS 60
                                                                                                                                     DIVMTQSPAIMSASPGEKVTITCSASSSV-----SYMHWFQQKPGTSPKLWLYSTSNLAS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Application US/08434000A
                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                            (619) 552-0159
                                                                                                                                                                                                                                                                                          DESCRIPTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 storage
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81.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            : including application
: described below:
08/367,395
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                                                                                                                                                                                                      3
                                                                                                                                                                                                                    Score 439.5; DB 2;
Pred. No. 7.4e-36;
                                                                                                                                                                                                                                                                                         Guy's 13 Kappa
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1.44
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                                                                                                                                                                                                                                      Length 105;
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Sequence 12, Applicati Patent No. 6303341 GENERAL INFORMATION:

Application US/09312157

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APPLICANT: ANDREW C. HIATT, JULIAN

K.-C. MA, THOMAS LEHNER

TITLE OF INVENTION: IMMUNOGLOBULINS CONTAINING PROTECTION

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DESCRIPTION: Guy's 13 Kappa; SEQUENCE DESCRIPTION: SEQ ID NO: 12: US-09-312-157-12
Search completed: November 21, 2005, 12:07:35
Job time: 43.2564 secs
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                                                                                                                                                                                                                                                                  Query Match 74.5%; Score 439.5; DB 2; Length 105; Best Local Similarity 81.9%; Pred. No. 7.4e-36; Matches 86; Conservative 3; Mismatches 11; Indels 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE LISTING INFORMATION FOR SEQ ID NO: 12: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM COMPATIBLE
COMPUTER: IBM COMPATIBLE
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: WOR'D Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/312,157
FILING DATE: 14-May-1999
CLASSIFICATION NUMBER: 08/434,000
APPLICATION NUMBER: 08/434,000
FILING DATE: *CUNCHOWN'>
APPLICATION NUMBER: 08/434,000
FILING DATE: *CUNCHOWN'>
APPLICATION NUMBER: 08/434,000
FILING DATE: *CUNCHOWN'>
ATTORNEY/AGENT INFORMATION:
NAME: GUISE, Jeffrey W.
REGISTRATION NUMBER: 34,613
REFERENCE/DOCKET NUMBER: 212/127
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 552-8400
TELEFAX: (619) 552-0159
TELEX: 67-351
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 19
CORRESPONDENCE ADDRESS:
ADDRESSE: Lyon & Lyon
STREET: 633 West, Pifth Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ZIP: 90071
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
                                                                                          56 GVPARFSGSGSGTSYSLTISRMEAEDAATYYCHORTSYPYTFGGG 100
                                                                                                                      61 GVPARFSGSGSGTSYSLTISRMEAEDAATYYCQQSNEDPYTFGSG 105
                                                                                                                                                                              CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 105 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
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                                                                                                                                                                                                                                                                           11; Indels 5;
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Minimum
Maximum
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No.
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Perfect score:
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Maximum Match 100%
Listing first 45 summaries
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            Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB seq
        $50.000 to the contract of the
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// cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*

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// cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*

// cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.pep:*

// cgn2_6/ptodata/1/pubpaa/US11_PUBCOMB.pep:*
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US-09-144-886-89
US-10-632-706-85
US-09-144-886-88
US-09-976-787-8
US-09-976-787-8
US-10-203-399B-8
US-10-203-399B-8
US-10-203-399B-8
US-10-778-910-8
US-10-805-198-27
US-10-805-198-27
US-10-805-198-27
US-10-805-198-28
US-09-96-787-24
US-09-96-787-24
US-09-96-787-24
US-09-96-787-24
US-09-976-787-24
US-09-985-198-23
US-10-805-198-23
US-10-805-198-29
US-09-96-787-29
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Sequence 86, Appl Sequence 89, Appl Sequence 88, Appl Sequence 88, Appl Sequence 89, Appl Sequence 27, Appl Sequence 28, Appl Sequence 29, Appl Sequence 29,
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US-09-144-886-89 (US-09-144-886-89) (Sequence 89, Application US/09144886) (Patent No. US20020155114A1) (GENERAL INFORMATION: (APPLICANT: Marks, James D (APPLICANT: Ameredorfer, Peter (TITLE OF INVENTION: Therapeutic Monoclonal Antibodies That Neutralize (TITLE OF INVENTION: Botulinum Neurotoxins (PILS REFERENCE: 2500.117USO) (CURRENT APPLICATION NUMBER: US/09/144,886)	Query Match Query Match Query Match  Best Local Similarity 100.0%; Pred. No. 3.3e-44; Best Local Similarity 100.0%; Pred. No. 3.3e-44;  Matches 112; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  Matches 112; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  Qy 1	US-10-632-706-86  US-10-632-706-86  Sequence 86, Application US/10632706  Publication No. US20040175385A1  GENERAL INFORMATION:  APPLICANT: AMERS, D.  APPLICANT: AMERS, D.  APPLICANT: AMERSDORFER, PETER  ITILE OF INVENTION: THERAPBUTIC MONOCLONAL ANTIBODIES THAT NEUTRALIZE BOTULINUM  TITLE OF INVENTION: NURDERCOXINS  FILE REFERENCE: 407T-895120US  FRIOR APPLICATION NUMBER: US/10/632,706  CURRENT APPLICATION NUMBER: US/0/400,721  FRIOR APPLICATION NUMBER: US 60/400,721  FRIOR FILING DATE: 1998-08-31  NUMBER OF SEQ ID NOS: 278  SOPTWARE: Patentin version 3.2  SEQ ID NO 86  LENGTH: 112  TYPE: PRT ORGANISM: Artificial FRATURE:  FRIOR ATTIFICATION: Single chain antibody  US-10-632-706-86	28 449.5 76.2 108 3 US-09-910-099-9 Sequence 9, Appli 29 449.5 76.2 235 3 US-09-910-099-17 Sequence 17, Appl 30 448.5 76.0 256 4 US-10-247-488-2 Sequence 2, Appli 31 448.5 76.0 258 4 US-10-247-488-4 Sequence 2, Appli 32 443.5 75.2 109 3 US-09-144-886-91 Sequence 91, Appli 33 442.5 75.0 108 4 US-10-803-622-240 Sequence 91, Appli 34 42.5 75.0 108 4 US-10-803-622-240 Sequence 240, Appli 34 42.5 74.5 105 5 US-10-781-989-12 Sequence 12, Appli 36 439.5 74.5 109 4 US-10-312-316-50 Sequence 12, Appli 39 438.5 74.5 109 4 US-10-312-316-50 Sequence 12, Appli 438.5 74.3 108 4 US-10-803-622-244 Sequence 244, Appli 438.5 74.3 108 4 US-10-803-653-244 Sequence 244, Appli 438.5 74.3 108 5 US-10-603-653-244 Sequence 244, Appli 438.5 74.3 244 5 US-10-609-671-2 Sequence 2, Appli 438.5 74.3 244 5 US-10-609-671-2 Sequence 2, Appli 44 437.5 74.2 110 4 US-10-803-622-237 Sequence 2, Appli 5 Sequence 2, Appli 5 Sequence 2, Appli 5 Sequence 2, Appli 6 Sequence 2, Appli 6 Sequence 2, Appli 6 Sequence 2, Appli 7 Sequence 2, Appli 8 Sequenc

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CURRENT APPLICATION NUMBER: US/10/632,706
CURRENT FILING DATE: 2003-08-01
PRIOR APPLICATION NUMBER: US 60/400,721
PRIOR FILING DATE: 2002-08-01
PRIOR APPLICATION NUMBER: US 09/144,806
PRIOR FILING DATE: 1998-08-31
NUMBER OF SEQ ID NOS: 278
SOFTWARE: Patentin version 3.2
SEQ ID NO 85
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GENERAL INFORMATION:
APPLICANT: Marks, James D
APPLICANT: Amersdorfer, Peter
TITLE OF INVENTION: Therapeutic Monoclonal Antibodies
TITLE OF INVENTION: Botulinum Neurotoxins
                                                                          Sequence 88, Application UP Atent No. US20020155114A1 GENERAL INFORMATION:
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NUMBER OF SEQ ID NOS: 98
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 89
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Best Local Similarity
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Best Local :
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APPLICANT: AMERSDORFER, PETER
TITLE OF INVENTION: THERAPEUTIC MONOCLONAL ANTIBODIES THAT NEUTRALIZE BOTULINUM
TITLE OF INVENTION: NEUROTOXINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILE REFERENCE: 407T-895120US
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ORGANISM: Artificial Sequence
PEATURE:
OTHER INFORMATION: Description of Artificial Sequence: BONT/A clone
OTHER INFORMATION: 3C3 region VL epitope 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT ORGANISM: Artificial FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OTHER INFORMATION: single chain antibody
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Pred. No. 3e+37;
1; Mismatches 4;
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Pred. No. 9.5e-42;
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GENERAL INFORMATION:
APPLICANT: Zhu, Zhenping
APPLICANT: Witte, Larry
ITILE OF INVENTION: Antibodies Specific to KD
FILE REFERENCE: 11245/46505
CURRENT APPLICATION NUMBER: US/09/976,787
CURRENT FILING DATE: 2001-10-12
PRIOR APPLICATION NUMBER: US 09/493,539
PRIOR APPLICATION NUMBER: US 60/117,726
PRIOR APPLICATION NUMBER: US 60/117,726
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Sequence 8, Application US/09865198
Patent No. US20020103345A1
GENERAL INFORMATION:
APPLICANT: Zhun, Zhenping
TITLE OF INVENTION: Bispecific Immunoglobulin-Like Antigen
TITLE OF INVENTION: Production
FILE REFERENCE: 11245/47102
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CURRENT APPLICATION NUMBER: US/09/144,886
CURRENT FILING DATE: 1998-08-31
NUMBER OF SEQ ID NOS: 98
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 88
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Best Local
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TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: PRT
ORGANISM: Mus musculus
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82.1%;
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Pred. No. 7.6e-36;
2; Mismatches 7;
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                                                                                                                                                                                                                                                                                                                                                                        Score 460.5; DB 3; Pred. No. 7.4e-33;
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CURRENT APPLICATION NUMBER: US/09/865,198

Binding Proteins and

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LENGTH: 108
TYPE: PRT
ORGANISM: Mouse
US-09-865-198-8
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Publication No. US20040242851A1
GENERAL INFORMATION:
APPLICANT: Zhu, Zhenping
TITLE OF INVENTION: Bispecific Antibodies That Bind to VEGF Receptors
FILE REFERENCE: 11245/48502
CURRENT APPLICATION NUMBER: US/10/482,630
CURRENT APPLICATION NUMBER: DO3-12-23
PRIOR APPLICATION NUMBER: PCT/US02/20332
PRIOR APPLICATION NUMBER: PCT/US02/20332
PRIOR APPLICATION NUMBER: US 60/301,299
PRIOR PILING DATE: 2001-06-26
PRIOR PILING DATE: 2001-06-26
NUMBER OF SEQ ID NOS: 137
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PRIOR APPLICATION NUMBER: US 60/206,749
PRIOR FILING DATE: 2000-05-24
NUMBER OF SEQ ID NOS: 34
SOFTWARE: WordPerfect 8.0 for Windows
SEQ ID NO 8
                                                                                                                                                                                 Sequence 8, Application US/10203399B Publication No. US20040248781A1 GENERAL INFORMATION:
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SEQ ID NO 8
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APPLICANT: Kerbel, Robert
TITLE OF INVENTION: Therapeutic Method for Reducing Angiogenesis
FILE REFERENCE: 11245/47002
CURRENT APPLICATION NUMBER: US/10/203,399B
CURRENT FILING DATE: 2002-08-08
PRIOR APPLICATION NUMBER: 601/78,791
PRIOR APPLICATION NUMBER: 601/78,791
PRIOR FILING DATE: 2000-01-28
PRIOR FILING DATE: 2000-03-31
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1 Similarity 82.1%;
92; Conservative
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82.1%;
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Pred. No. 7.4e-33;
4; Mismatches 11;
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Sequence 8, Application US/10809387
Publication No. US20050214860A1
GENERAL INFORMATION:

APPLICANT: Zhu, Zhenping
APPLICANT: Witte, Larry
TITLE OF INVENTION: Antibodies Specific to KDR
FILE REFERENCE: 11245/46506
CURRENT APPLICATION NUMBER: US/10/809,387
CURRENT FILING DATE: 2004-03-23
PRIOR APPLICATION NUMBER: 09/976,787
PRIOR APPLICATION NUMBER: 09/976,787
PRIOR APPLICATION NUMBER: 09/493,539

and Uses Thereof

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; LENGTH: 108
; TYPE: PRT
; ORGANISM: Mouse
US-10-778-910-8
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US-10-778-910-8
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US-10-809-387-8
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PRIOR PILING DATE: 2001-01-29
NUMBER OF SEQ ID NOS: 16
SOFTWARE: PatentIn Ver. 2.1
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TITLE OF INVENTION: Bispecific Immunoglobulin-Like Antigen Binding Proteins and
TITLE OF INVENTION: Production
FILE REFERENCE: 11245/47102
CURRENT FILING DATE: 2004-02-13
PRIOR APPLICATION NUMBER: US/10/778,910
CURRENT FILING DATE: 2004-02-13
PRIOR APPLICATION NUMBER: US/09/865,198
PRIOR FILING DATE: 2001-05-24
PRIOR APPLICATION NUMBER: US 60/206,749
PRIOR APPLICATION DATE: 2000-05-24
NUMBER OF SEQ ID NOS: 34
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TYPE: PRT
ORGANISM: Mouse
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                                                                                                                    61 GVPARFSGSGSGTSYSLTISRMEAEDAATYYCQQSNEDPYTFGSGDQAGNKR 112
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82.1%;
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82.1%;
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Pred. No. 7.4e-33;
4; Mismatches 11;
                                                                                                                                                                                                                                         4; Mismatches
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Pred. No. 7.4e-33;
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SEQ ID NO 8

LENGTH: 108

PRIOR FILING DATE: 1999-01-29 NUMBER OF SEQ ID NOS: 40 SOFTWARE: WordPerfect 8.0 for Windows

PRIOR FILING DATE:

FILING DATE: 2000-01-28
APPLICATION NUMBER: 60/117,726

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; ORGANISM: Mouse
US-09-976-787-28
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APPLICANT: Mitte, Larry
APPLICANT: Mitte, Larry
ITILE OF INVENTION: Antibodies Specific to KD:
FILE REFERENCE: 11245/46505
CURRENT APPLICATION NUMBER: US/09/976,787
CURRENT FILING DATE: 2001-10-12
PRIOR APPLICATION NUMBER: US 09/493,539
PRIOR PILING DATE: 2000-01-28
PRIOR APPLICATION NUMBER: US 60/117,726
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US-09-976-787-28
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Sequence 27, Application US/09865198
Patent No. US20020103345A1
GENERAL INFORMATION:
APPLICANT: Zhu, Zhenping
TITLE OF INVENTION: Bispecific Immunoglobulin-Like Antigen Binding Proteins and Methorities of INVENTION: Production
FILE REFERENCE: 11245/47102
CURRENT APPLICATION NUMBER: US/09/865,198
CURRENT FILING DATE: 2001-05-24
PRIOR APPLICATION NUMBER: US 60/206,749
PRIOR FILING DATE: 2000-05-24
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Patent No. US20020064528A1
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Best Local
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Best Local Similarity 82.1%;
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NUMBER OF SEQ ID NOS:
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Local Similarity 82.1%;
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Pred. No. 7.4e-33;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 460.5; DB 3
Pred. No. 1.7e-32;
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SOFTWARE: WordPerfect

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; TYPE: PRT
; ÖRGANISM: Mouse
US-09-865-198-27
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Best Local S
Matches 92
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LENGTH: 240
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SOPTWARE: WordPerfect 8.0
SEQ ID NO 27
                                                                                                                                                                                                                                                             Sequence 27, Application US/10778910 Publication No. US20040259156A1 GENERAL INFORMATION:
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TITLE OF INVENTION: Bispecific Antibodies That Bind to VEGF Receptors
FILE REFERENCE: 11245/48502
CURRENT APPLICATION NUMBER: US/10/482,630
CURRENT FILING DATE: 2003-12-23
PRIOR APPLICATION NUMBER: PCT/US02/20332
PRIOR APPLICATION NUMBER: PCT/US02/20332
PRIOR TILING DATE: 2002-06-26
PRIOR TILING DATE: 2002-06-26
TITLE OF INVENTION: Bispecific Immunoglobulin-Like Antigen Binding Proteins TITLE OF INVENTION: Production FILE REFERENCE: 11245/47102
CURRENT APPLICATION NUMBER: US/10/778,910
CURRENT FILING DATE: 2004-02-13
PRIOR APPLICATION NUMBER: US/09/865,198
PRIOR PILING DATE: 2000-05-24
NUMBER OF SEQ ID NOS: 34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: US 60/301,299
PRIOR FILING DATE: 2001-06-26
NUMBER OF SEQ ID NOS: 137
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: WordPerfect 8.0 for Windows
                                                                                                                                                                                                                                          APPLICANT: Zhu, Zhenping
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82.1%; Pred. No. 1.7e-32;
Live 4; Mismatches 11;
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Sequence 28, Application US/10809387

Publication No. US20050214860A1

GENERAL INFORMATION:

APPLICANT: Zhu, Zhenping

APPLICANT: Witte, Larry

ITILE OF INVENTION: Antibodies Specific to KDR and Uses Thereof

FILE REFERENCE: 11245/46506

CURRENT APPLICATION NUMBER: US/10/809,387

CURRENT FILING DATE: 2004-03-23

PRIOR APPLICATION NUMBER: 09/976,787

PRIOR FILING DATE: 2001-10-12

PRIOR FILING DATE: 2001-0-128

PRIOR APPLICATION NUMBER: 60/117,726

PRIOR APPLICATION NUMBER: 60/117,726

PRIOR APPLICATION NUMBER: 60/117,726

PRIOR APPLICATION NUMBER: 60/117,726

PRIOR FILING DATE: 1999-01-29

NUMBER OF SEQ ID NOS: 40

SOFTWARE: WordPerfect 8.0 for Windows

SEQ ID NO 28

LENGTH:: 240

TYPE: PAT

ORGANISM: Mouse

US-10-809-387-28
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; LENGTH: 240
; TYPE: PRT
; ORGANISM: Mouse
US-10-778-910-27
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Search completed: November 21, 2005, 12:33:33 Job time: 139.667 secs
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Scoring table:
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Maximum Match 100%
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Maximum DB seq length: 2000000000
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     Pr&A. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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1: /cgn2 6/ptodata/2/pubpaa/US10 NEW PUB.pep:*
2: /cgn2_6/ptodata/2/pubpaa/US06 NEW PUB.pep:*
3: /cgn2_6/ptodata/2/pubpaa/US07 NEW PUB.pep:*
4: /cgn2_6/ptodata/2/pubpaa/US08 NEW PUB.pep:*
5: /cgn2_6/ptodata/2/pubpaa/US09 NEW PUB.pep:*
6: /cgn2_6/ptodata/2/pubpaa/US1_NEW PUB.pep:*
7: /cgn2_6/ptodata/2/pubpaa/US1_NEW PUB.pep:*
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Match
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Copyright (c) 1993 - 2005 Compugen Ltd.
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US-11-177-978-1
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US-11-174-186-8
US-11-174-186-8
US-11-174-186-3
US-11-174-186-16
US-10-512-184-36
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US-11-174-186-14
US-11-174-186-17
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RESULT 1 US-10-495-664-3 US-10-495-664 ; Sequence 3, Application US/1049564 ; Publication No. US20050244416A1 [GENERAL IMPORMATION: IGENERAL IMPORMATION: IGENERAL IMPORMATION: REPLICATION UNDER ITTIL OF INVERTION: PILIS REPERENCE .004218-0001 PILIS REPERENCE .004289-0001 PILIS REPERENCE .004289-001 PI	26 339 57.5 129 1 US-10-721-763-23 Sequence 23, Appli 27 338 57.3 110 1 US-10-648-816-1 Sequence 5, Appli 28 334.5 56.9 110 1 US-10-648-816-1 Sequence 1, Appli 39 34.5 56.7 213 1 US-10-721-763-27 Sequence 27, Appli 39 35.6 4 131 1 US-10-512-184-29 Sequence 27, Appli 39 39.5 55.8 106 7 US-11-174-186-1 Sequence 29, Appli 39 39.5 55.8 213 7 US-11-174-186-5 Sequence 5, Appli 39 39.5 55.8 213 7 US-11-174-280-8 Sequence 64, Appli 39 317 55.4 96 7 US-11-144-280-42 Sequence 67, Appli 39 317 53.7 126 1 US-10-839-799-65 Sequence 61, Appli 39 317 53.7 126 1 US-10-839-799-65 Sequence 65, Appli 39 317 53.7 126 1 US-10-839-799-65 Sequence 65, Appli 39 317 53.7 126 1 US-10-839-799-65 Sequence 65, Appli 39 317 53.6 126 1 US-10-839-799-65 Sequence 65, Appli 39 318 53.4 107 1 US-10-839-799-65 Sequence 65, Appli 39 318 53.4 107 1 US-10-839-799-65 Sequence 65, Appli 39 318 53.2 236 7 US-11-144-248-51 Sequence 65, Appli 39 318 53.2 236 7 US-11-144-248-51 Sequence 66, Appli 39 318 53.2 236 7 US-11-144-248-51 Sequence 67, Appli 39 318 53.2 236 7 US-11-144-248-51 Sequence 67, Appli 39 318 53.1 251 1 US-10-512-184-67 Sequence 67, Appli 39 318 53.1 320 1 US-10-512-184-67 Sequence 67, Appli 39 318 53.1 320 1 US-10-512-184-67 Sequence 67, Appli 39 318 53.1 320 1 US-10-512-184-67 Sequence 67, Appli 39 318 53.1 320 1 US-10-512-184-67 Sequence 67, Appli 39 318 53.1 320 1 US-10-512-184-67 Sequence 67, Appli 39 318 53.1 320 1 US-10-512-184-67 Sequence 67, Appli 39 318 53.1 320 1 US-10-512-184-67 Sequence 67, Appli 39 318 53.1 320 1 US-10-512-184-67 Sequence 67, Appli 30 318 53.1 320 1 US-10-512-184-67 Sequence 67, Appli 30 318 53.1 320 1 US-10-512-184-67 Sequence 67, Appli 30 318 53.1 320 1 US-10-512-184-67 Sequence 67, Appli 30 318 53.1 320 1 US-10-512-184-67 Sequence 67, Appli 30 318 53.1 320 1 US-10-512-184-67 Sequence 67, Appli 30 318 53.1 320 1 US-10-512-184-67 Sequence 67, Appli 30 318 53.1 320 1 US-10-512-184-67 Sequence 67, Appli 30 318 53.1 320 1 US-10-512-184-67 Sequence 67, Appli 30 318 53.1 320 1 US-10-512-184-67 Sequence 67, Appli
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US-11-077-978-1
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APPLICANT: Yazaki, Paul J.
APPLICANT: Shively, John B.
APPLICANT: Shively, John B.
APPLICANT: Raubitschek, Andrew A.
APPLICANT: Raubitschek, Andrew A.
APPLICANT: Wu. Anna M.
APPLICANT: Wu. Anna M.
TITLE OF INVENTION: A Humanized Anti-CEA T84.66 Antibody and Uses Thereof FILE REFERENCE: 54435.8012.US01
CURRENT APPLICATION NUMBER: US/11/077,978
CURRENT FILING DATE: 2005-03-11
NUMBER OF SEQ ID NOS: 27
SOFTWARE: Patentin version 3.2
SEQ ID NO 1
LENGTH: 111
Sequence 8, Application US/11174186
Publication No. US20050244418A1
GENERAL INFORMATION:
APPLICANT: Gillies, Stephen
APPLICANT: Lo, Kin-Ming
APPLICANT: Qian, Xiugi
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Best Local Similarity
Matches 71; Conserv
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CURRENT FILING DATE: 2005-07-01
PRIOR APPLICATION NUMBER: US 60/288,564
PRIOR FILING DATE: 2001-05-03
NUMBER OF SEQ ID NOS: 42
SOFTWARE: PATENTIN VERSION 3.3
SEQ ID NO 1
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TYPE: PRT
ORGANISM: Artificial
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Pred. No. 3.7e-21;
5; Mismatches 15
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Pred. No. 1e-20;
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US-11-174-186-9
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Publication No. US20050244418A1
GENERAL INFORMATION:
APPLICANT: Gillies, Stephen
APPLICANT: Qian, Xiugi
APPLICANT: Qian, Xiugi
TITLE OF INVENTION: Recombinant Tumor Specific Antibody And Use Thereof
FILE REFERENCE: LEX-019
FULGRENT APPLICATION NUMBER: US/11/174,186
CURRENT FILING DATE: 2001-05-03
PRIOR APPLICATION NUMBER: US 60/288,564
PRIOR FILING DATE: 2001-05-03
NUMBER OF SEQ ID NOS: 42
GOTTMAND: Deferition of the control of th
Sequence 9, Application US/11174186
Publication No. US20050244418A1
GENERAL INFORMATION:
APPLICANT: Gillies, Stephen
APPLICANT: Lo, Kin-Ming
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SEQ ID NO 42
LENGTH: 213
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CURRENT APPLICATION NUMBER: US/11/174,186
CURRENT FILING DATE: 2005-07-01
PRIOR APPLICATION NUMBER: US 60/288,564
PRIOR FILING DATE: 2001-05-03
NUMBER OF SEQ ID NOS: 42
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Similarity 68.8%; Pred. No. 2.3
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                                                                                                                                             RESULT 8
US-10-512-184-50
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CURRENT APPLICATION NUMBER: US/10/512,184
CURRENT FILING DATE: 2004-10-22
NUMBER OF SEQ ID NOS: 72
SOFTWARE: Patentin Ver. 2.1,
SEQ ID NO 36
LENGTH: 248
TURE TO THE TITLE OF THE T
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TITLE OF INVENTION: Recombinant Tumor Specific Antibody And Use Thereof
FILE REFERENCE: LEX-019
CURRENT APPLICATION NUMBER: US/11/174,186
CURRENT FILING DATE: 2005-07-01
PRIOR APPLICATION NUMBER: US 60/288,564
PRIOR APPLICATION NUMBER: US 60/288,564
RIOR FILING DATE: 2001-05-03
NUMBER OF SEQ ID NOS: 42
SOFTWARE: Patentin version 3.3
SEQ ID NO 9
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Best Local
                           Sequence 50, Application US/10512184
Publication No. US20050244901A1
GENERAL INFORMATION:
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Best Local :
APPLICANT: Fraunhofer Gesellschaft zur F"rderung der angewandten Forschung e.V.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OTHER INFORMATION: Description of Artificial Sequence: scFv SS2 with OTHER INFORMATION: specificity against Sclerotinia sclerotiorum; OTHER INFORMATION: originates from Mus musculus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Artificial Sequence FEATURE:
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Local Similarity 71.4%;
les 75; Conservation
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Pred. No. 2.6e-20;
8; Mismatches 17
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LENGTH: 106
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Best Local &
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                                                                                                                                                                    Matches
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APPLICANT: Gillies, Kiumi
APPLICANT: Oin, Kiumi
APPLICANT: Oin, Kiumi
APPLICANT: Oin, Kiumi
TITLE OF INVENTION: Recombinant Tumor Specific Antibody And Use Thereof
TITLE REFERENCE: LEX-019
CURRENT APPLICATION NUMBER: US/11/174,186
CURRENT FILING DATE: 2005-07-01
PRIOR APPLICATION NUMBER: US 60/288,564
PRIOR PILING DATE: 2001-05-03
NUMBER OF SEQ ID NOS: 42
SOFTWARE: Fatentin version 3.3
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ORGANISM: Artificial
FEATURE:
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ORGANISM: Artificial Sequence
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57 FPARFSGSGSGTSYTLTINSLEAEDAATYYCHORSGYPYTFGGG 100
                                     62 VPARFSGSGSGTSYSLTISRMEAEDAATYYCQQSNEDPYTFGSG 105
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                                                                                                                                                                                      62.6%; Score 369.5; DB 7; 68.3%; Pred. No. 1.6e-19;
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/ Sequence 16, Application US/11174186
/ Publication No. US20050244418A1
/ GENERAL INFORMATION:

RESULT 10 US-11-174-186-16

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RESULT 12
US-11-174-186-10
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APPLICANT: Qian, Xiugi

TITLE OF INVENTION: Recombinant Tumor Specific Antibody And Use Thereof

FILE REFERENCE: LEX-019

CURRENT APPLICATION NUMBER: US/11/174,186

CURRENT FILING DATE: 2005-07-01

PRIOR APPLICATION NUMBER: US 60/288,564

PRIOR PILING DATE: 2001-05-03

NUMBER OF ERQ ID NOS: 42

SOFTWARE: Patentin version 3.3

SEQ ID NO 16
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SEQ ID NO 4
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Best Local &
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APPLICANT: Wu, Anna M.
APPLICANT: Wu, Anna M.
TITLE OP INVENTION: A Humanized Anti-CEA T84.66 Antibody and Uses Thereof
FILE REFERENCE: 54435,8012.US01
CURRENT APPLICATION NUMBER: US/11/077,978
CURRENT FILING DATE: 2005-03-11
NUMBER OF SEQ ID DATE: 2005-03-12
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                                                                                                                                                                                                                                                                                                                                                 PUBLICATION INPORMATION:
DATABASE ACCESSION NUMBER: 1MOEA
DATABASE ENTRY DATE: 2002-09-09
RELEVANT RESIDUES: (1)..(111)
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APPLICANT: Sherman, Mark
APPLICANT: Shively, John
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ORGANISM: Artificial
FEATURE:
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Shively, John E.
Raubitschek, Andrew A.
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61.9%; Pred. No. 7.8e-19;
tive 15; Mismatches 25;
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Pred. No. 1.8e-19;
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US-11-174-186-14
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Publication No. US20050244418A1 
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 10, Application US/11174186
Publication No. US20050244418A1
GENERAL INFORMATION:
APPLICANT: G1116e, Stephen
APPLICANT: Lo, Kin-Ming
APPLICANT: Qian, Xiugi
TITLE OF INVENTION: Recombinant Tumor Specific Antibody And Use Thereof
FILE REFERENCE: LEX-019
CURRENT FILING DATE: 2005-07-01
PRIOR APPLICATION NUMBER: US 60/288,564
PRIOR FILING DATE: 2001-05-03
                                                                                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn version 3.3 SEQ ID NO 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOPTWARE: PatentIn version 3.3
SEQ ID NO 10
LENGTH: 106
                                                                                                                                                     Matches
                                                                                                                                                                    Query Match
Best Local Similarity
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APPLICANT: Lo, Kin-Ming
APPLICANT: Qian, Xingi
TITLE OF INVENTION: Recombinant Tumor Specific Antibody And Use Thereof
FILE REPERRNCE: LEX-019
CURRENT APPLICATION NUMBER: US/11/174,186
CURRENT FILING DATE: 2005-07-01
                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: US 60/288,564 PRIOR FILING DATE: 2001-05-03 NUMBER OF SEQ ID NOS: 42
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                                                                                                                                                                                                                                                                                  ORGANISM: Artificial
                                                                                                                                                                                                                                                                                                        TYPE: PRT
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57
                   62 VPARFSGSGSGTSYSLTISRMEAEDAATYYCQQSNEDPYTFGSG 105
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FPARFSGSGSGTSYTLTINSLEAEDAATYYCHQRSGYPYTFGGG 100
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                                                                       ILLTOSPASLAVSPGORATITCSASSSV----SYMLWYQQXPGQPPKPWIFDTSNLASG 56
                                                                                                       ISLTQSPAIMSASPGEKVTTTCRASSSVDSYGHSFMQWFQQKPGTSPKLWIYSTSNLASG 61
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                                                                                                                                                   Conservative
                                                                                                                                                                    60.8%; Score 358.5; DB 7 66.3%; Pred. No. 8.1e-19;
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66.3%; Pred. No. 8.1e-19;
:ive 11; Mismatches 19
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CURRENT APPLICATION NUMBER: US/11/174,186
CURRENT FILING DATE: 2005-07-01
PRIOR APPLICATION NUMBER: US 60/288,564
PRIOR FILING DATE: 2001-05-03
NUMBER OF SEQ ID NOS: 42
SOFTWARE: PATENTIN VERSION 3.3
SEQ ID NO 13
LENGTH: 106
TYPE: PRT
ORGANISM: Artificial
PERTURE:
COMMENT APPLICATION.
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US-11-174-186-13
US-11-174-186-13
Sequence 13, Application US/11174186
; Sequence 10, Application US-11174186
; Publication No. US20050244418A1
; GENERAL INFORMATION:
; APPLICANT: Gillies, Stephen
; APPLICANT: Lo, Kin-Ming
; APPLICANT: Lo, Kin-Ming
; APPLICANT: Olan, Xiugi
; TITLE OF INVENTION: Recombinant Tumor Specific Antibody And Use Thereof
; FILE REFERENCE: LEX-019
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; TYPE: PRT
; ORGANISM: Artificial
; PEATURE:
; PEATURE:
; OTHER INFORMATION: KS de-immunized VK2
US-11-174-186-12
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APPLICANT: Gillies, Stephen
APPLICANT: Gillies, Stephen
APPLICANT: Uo, Kin-Ming
APPLICANT: Qian, Xiusi
TITLE OF INVENTION: Recombinant Tumor Specific Antibody And Use Thereof
FILE REFERENCE: LEX-019
CURRENT APPLICATION NUMBER: US/11/174,186
CURRENT FILING DATE: 2005-07-01
PRIOR APPLICATION NUMBER: US 60/288,564
PRIOR FILING DATE: 2001-05-03
NUMBER OF SEQ ID NOS: 42
SOPTWARE: Patentin version 3.3
SEQ ID NO 12
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                                                                                                                                                                       Query Match 60.3%; Score 355.5; DB 7; Length 106; Best Local Similarity 65.4%; Pred. No. 1.3e-18; Matches 68; Conservative 12; Mismatches 19; Indels 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        y Match 60.4%; Score 356.5; DB 7; Length 106; Local Similarity 65.4%; Pred. No. 1.1e-18; hes 68; Conservative 12; Mismatches 19; Indels 5
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                                                                                       ILITQSPASLAVSPGQRATITCSASSSV-----SYMLWYQQKPGQPPKPWIFDTSNLASG 56
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seq length: 2000000000
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                                                                                                                                            A_Geneseq_21:*
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Copyright (c) 1993 - 2005 Compugen Ltd.
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No.	Score	54	Length	B	! !	Description
_	553	100.0	107	8	ADR38685	ouse
N	518	•	109	8	ADR38686	Adr38686 Mouse lig
ω	516	93.3	107	8	ADR38673	
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6	515	93.1	210	8	ADU86565	Adu86565 Immunoglo
7	515	93.1	210	8	ADU86571	Adu86571 Immunoglo
œ	510	92.2	108	8	ADT74993	Light
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10	506	91.5	244	N	AAR79867	
Ħ	505		242	N	AAR79871	
12	502	90.8	106	v	ABG32682	Murin
13	502		106	v	AAE27810	
14	502	•	128	N	AAR27050	
15	501	90.6	106	4	AAB69673	Murine
16	501	90.6	106	8	ADO47720	Ado47720 Mouse mik
17	501	90.6	128	4	AAB69655	Murine
18	501	90.6	128	8	ADO47747	
19	499	90.2	242	N	AAR79870	
20	499		242	N	AAR79872	
21	498.5	90.1	108	9	ADW86127	
22	498.5	90.1	214	8	ADU86519	
23	497	89.9	106	8	AD016879	Ado16879 Murine an
24	496	89.7	246	N	AAR79869	Aar79869 Anti-EGFR

The invention describes an isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a specific clone where (I) binds to and neutralises botulinum neurotoxin type A (BoNT/A). An isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a clone chosen from clone 25, C25, C39, 1C6, 3D12, B4, 1F3, huC25, A11, A22, WR1(V), WR1(T), 3-1, 3-8, 3-10 and ING1, where (I) binds to and neutralizes botulinum neurotoxin type A (BoNT/A). Also described are: a polypeptide (II) comprising BoNT/A neutralising epitope having an epitope that is

New isolated antibody that neutralizes botulinum neurotoxin type A, useful for diagnosing botulism or for treating pathologies associated with botulinum neurotoxin poisoning.

Example 1; SEQ ID NO 87; 110pp; English.

WPI; 2004-652009/63.

Marks JD,

Amersdorfer P;

45	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30	29	28	27	26	25
484	484	484	485	485	486	486	486	486	488	488	488	488	488	488	488	488	488	489	489	491.5
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ADH77287	ADW64803	ADW64805	AAR21301	ADW80852	ADX69520	ADZ21542	AAG66521	ADZ21534	ADZ21537	AAE33333	ADZ21541	AAR79873	AAR79868	AAR12327	ADZ21532	ADZ21524	AAE33326	AEA29819	AAR21293	AAY93891
Adh77287					_		•	-		. ~			_					_	-	٠.
Yeast kil		Murine 81	Murine VL	Anti-CD3	Mouse ICA	Anti-Muc-	Mouse ant	Anti-Muc-	Anti-Muc-	12E ancio	Anti-Muc-	Anti-Burk	ANCI-EGER	Light (ka	Anci-Muc-	Anti-Muc-	128 aucio	Mouse nig	Murine vi	The kappa

## ALIGNMENTS

## RRESULT 1 ADR38621 ID ADR383 XX ADR33 XX ADR3 AC ADR3 XX ADR3 XX MOUS XX MOUS XX MOUS XX HOUS XX US20 XX I19h XX I19h XX I20 PN US20 XX I21 XX I21 XX I22 XX I23 XX I23 XX I24 XX I25 XX 31-AUG-1998; 98US-00144886. 01-AUG-2002; 2002US-0400721P. 01-AUG-2003; 2003US-00632706. 09-SEP-2004. antibacterial; antibody; botulinum neurotoxin type A; BONT/A; BONT/A neutralising epitope; anti-botulinum neurotoxin antibody; toxin neutralisation; botulinum neurotoxin poisoning; mouse; light chain variable region; single chain antibody; scFv. ADR38685 standard; peptide; 107 AA. US2004175385-A1. Mus sp Mouse light 02-DEC-2004 ADR38685; (REGC ) UNIV CALIFORNIA. chain variable region scFv seqid 87 (first entry)

specifically bound by an antibody expressed by clones as mentioned in (I), producing (I), and a composition (III) comprising several antibotuilnum neurocoxin antibodies, where each antibody is specific for a different epitope of a botulinum neurocoxin and the combination of

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ADR38686
ID ADR3
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Best Local S
Matches 107
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                                                                                                                                          useful for arayivering --- with botulinum neurotoxin poisoning.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    antibacterial; antibody; botulinum neurotoxin type A; BONT/A; BONT/A neutralising epitope; anti-botulinum neurotoxin antibody; toxin neurralisation; botulinum neurotoxin poisoning; mouse; light chain variable region; single chain antibody; scFv.
                                                                                                       Example 1; SEQ ID NO 88; 110pp; English.
                                                                                                                                                                                                        New isolated antibody that neutralizes botulinum neurotoxin type A, useful for diagnosing botulism or for treating pathologies associated
                                                                                                                                                                                                                                                                                                             WPI; 2004-652009/63.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               31-AUG-1998; 98US-00144886.
01-AUG-2002; 2002US-0400721P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FSGSGSGTSYSLTISRMEAEDAATYYCQQWSSNPLTFGSGTKLELKR 107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     standard; peptide; 109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DTELTQSPALMSASPGEKVTTTCSASSSVSYMYMYQQKPGSSPRLLIYDTSNLASGVPVR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PSGSGSGTSYSLTISRMEAEDAATYYCOOWSSNPLTFGSGTKLELKR 107
                                                                                                                                                                                                                                                                                                                                                                               Amersdorfer P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 107;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          60
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CC clone where (I) binds to and neutralises botulinum neurotoxin type A CC (BoNT/A). An isolated antibody (I) that specifically binds to an epitope CC specifically bound by an antibody expressed by a clone chosen from clone CS25, C25, C39, 1C6, 3D12, B4, 1F3, huC25, kT1, kT2, kT1(V), kT1(T), 3-1, CC 3-8, 3-10 and ING1, where (I) binds to ahd neutralizes botulinum CC neurotoxin type A (BONT/A). Also described are: a polypeptide (II) CC producing (I); and a composition (III) comprising source and in (I) comprising source and in (I) continum neurotoxin antibodies, where each antibody is specific for a clifferent epitope of a botulinum neurotoxin and the combination of CC antibodies shows greater toxin neutralisation than the single antibodies in surplus. The following are disclosed: a pharmaceutical composition CC comprising (I); and a kit comprising (I). (I) is useful for neutralising BONT/A antibody and for neutralising a botulinum neurotoxin which CC involves contacting neurotoxin with (I) in surplus, where each of (I) is specific for a different epitope of the botulinum neurotoxin and the combination of antibodies shows greater toxin neutralisation than the specific for a different epitope of the botulinum neurotoxin and the combination of antibodies shows greater toxin neutralisation than the single antibodies in surplus. (I) is useful for neutralisation of corporate antibodies shows greater toxin neutralisation than the single antibodies in surplus. (I) is useful for diagnosing the botulism corporate antibodies associated with botulinum neurotoxin and the combination of antibodies associated with botulinum neurotoxin corporates associated with botulinum neurotoxin acid sequence of a mouse light chain variable fragment anti-botulinum toxin care.
Sequence 109 AA;
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Query Match
Best Local S
Matches 103
                                                                                                                                       103;
61
                            59
                                                                  <u>, .</u>
                                                                                        1 DTELTQSPAIMSASPGEKVTTTCSASSSV--SYMYWYQQKPGSSPRLLIYDTSNLASGVP
                                                                                                                                                     Similarity
 VRFSGSGSGTSYSLTISRMEAEDAATYYCQQWSSNPLTFGSGTKLELKR
                   VRFSGSGSGTSYSLTISRMEAEDAATYYCQQWSSNPLTFGSGTKLELKR
                                                                DIELTQSPAIMSASPGEKVTTTCRASSSVSS6YLGWYQQKPGSSPRLLIYDTSNLASGVP 60
                                                                                                                                   93.7%;
ilarity 94.5%;
Conservative
                                                                                                                                   Score 518; DB 8; Length 109; Pred. No. 9.3e-32; Indels 1; Mismatches 3; Indels
 109
                                107
                                                                                                                                     <u>ب</u>
                                                                                                                                   Gaps
                                                                                                    58
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닭 ঠ

RESULT 3 ADR38673 THE SECOND PART OF THE SECOND PA Mouse light chain variable region scFv seqid 75. ADR38673 standard; peptide; 107 (first entry)

antibacterial; antibody; botulinum neurotoxin type A; BoNT/A; BoNT/A neutralising epitope; anti-botulihum neurotoxin antibody; toxin neutralisation; botulinum neurotoxin poisoning; mouse; 31-AUG-1998; 98US-00144886. 01-AUG-2002; 2002US-0400721P. 01-AUG-2003; 2003US-00632706. 09-SEP-2004. US2004175385-A1 de enw light chain variable region;

The invention describes an isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a specific

WPI; 2004-652009/63.

(REGC ) UNIV CALIFORNIA.

ğ,

Amersdorfer

New isolated antibody that neutralizes botulinum neurotoxin type

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RESULT 4
ABB75616
ID ABB7
XX ABB757
XX ABB7
XX ABB7
XX ABB7
XX ABB7
XX AOR
DT 10-J
DX AOR
DX 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention describes an isolated antibody (I) that specifically binds colone where (I) binds to and neutralises botulinum neurotoxin type A (BoNT/A). An isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a clone chosen from clone (C (BoNT/A). An isolated antibody (I) that specifically binds to an epitope expecifically bound by an antibody expressed by a clone chosen from clone (C 3-8, 3-10 and ING1, Where (I) binds to and neutralizes botulinum (C comprising BoNT/A neutralising epitope having an epitope that is groducing (I); and a composition (II) comprising several anticolor (I); producing (I); and a composition (II) comprising several anticolor (I); producing (I); and a composition (II) comprising several anticolor (I); and a present each antibody is specific for a different epitope of a botulinum neurotoxin and the combination of comprising (I); and a kit comprising (I) is useful for neutralising (I) and a kit comprising (I) is useful for neutralising (I) is useful for neutralising (I) and a kit comprising (I) is useful for neutralising (I) is useful for a different epitope of the botulinum neurotoxin which (I) is specific for a different epitope of the botulinum neurotoxin and the combination of antibodies shows greater toxin neutralisation than the single antibodies in surplus. (I) is useful for diagnosing the botulism (I) exhibits specificity and affinity towards BONT/A. (I) enables rapid detection or diagnosis of botulism. This is the amino acid composition of a mouse light chain variable fragment anti-botulinum toxin (I) serve.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               밁
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches 101;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    useful for diagnosing botulism or for treating pathologies associated with botulinum neurotoxin poisoning.
                                                                                                                                                                                                                                                                                                                                             RANK ligand; RANK-L; monoclonal antibody; mAb; 19H22; antibody; complementarity determining region; CDR; rheumatoid arthritis; osteoporosis; bone cancer; metastasis; osteolysis; osteoarthritis; psoriasis; inflammatory bowel disease; multiple sclerosis; diabetes; cytostatic; antiarthritic; antiinflammatory; osteopathic; antidiabetic; antirheumatic; immunosuppressive; antipsoriatic; neuroprotective;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 107 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 1; SEQ ID NO 75; 110pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Anti-RANK ligand mAb 19H22 light chain variable region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           10-JUN-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABB75616 standard; protein;
                                    Region
                                                                                                                                       Region
                                                                                                                                                                                                                                                 Mus musculus.
                                                                                                                                                                                                                                                                                                                       Light
                                                                                                                                                                                                                                                                                                                       chain; VL; mouse.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 DTBLTQSPAIMSASPGEKVTTTCSASSSVSYMYWYQQKPGSSPRLLIYDTSNLASGVPVR 60
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FSGSGSGTSYSLTISRMEAEDAATYYCQQWSSNPLTFGSGTKLELKR 107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PSGSGSGTSYSLTISRMEAEDAATYYCQQWSSYPLTPGAGTKLELKR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                          /label= CDR1
                                                                                                                                                                        Location/Qualifiers
       /label= CDR2
                                                                     note= "complementarity determining region"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  93.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          129 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 516; DB 8; Length 107; Pred. No. 1.3e-31; Mismatches 5; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   107
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              60
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21-APR-2005 ADW86126; ADW86126

(first entry)

standard;

protein; 107 AA

63

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RESULT 5
ADW86126
ID ADW8
XX
AC ADW8
XX
AC ADW8
XX
XX
XX
XX
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                                                                                                                                                                                                                                                                                                                                                                                                                            Cc claimed anti-human RANK ligand (RANK-L) murine monoclonal antibody (mab) cc laimed anti-human RANK ligand (RANK-L) murine monoclonal antibody (mab) cobtained by immunising CB6 f1 mice with human RANK-L protein, and fusing spleen cells with X61-A98.63 myeloma cells. 19H22 mb is specific for cc man RANK-L, having a binding affinity characterised by a dissociation constant of about 10 power -10 M. RANK-L is a member of the tumour constant of about 10 power -10 M. RANK-L is a member of the tumour constant of about 10 power -10 M. RANK-L is a member of the tumour chimeric and humanised 19H22 antibodies. The invention provides altered, chimeric and humanised 19H22 antibodies, including Fab or F(ab')2 cc fragments, antibodies comprising the light and heavy chain variable cregions of 19H22, and antibodies comprising the complementarity conditions of 19H22. These are used in methods for the constant dark induced osteoperic diseases, including concer, constants induced osteoperis, metastatic and primary bone cancer, constants induced osteoperis, metastatic and primary bone cancer, constants induced osteoperis, metastatic, and immune diseases concluding psoriasis, insulin dependent diabetes, inflammatory bowel concernation of the antibodies, using isolated mucleic acids constants conditions associated with Thi T-cell activity or osteoclast condevelopment and activation, in particular those listed above
                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A mouse monoclonal antibody 19H22 for treating or preventing osteopenic disease including rheumatoid arthritis and osteoprosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sweet RW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21-AUG-2000; 2000US-0226524P.
07-SEP-2000; 2000US-0230639P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         28-FEB-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO200215846-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present sequence is that of the light chain variable region of claimed anti-human RANK ligand (RANK-L) murine monoclonal antibody
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 6; Page 7; 45pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  N-PSDB; ABL53530.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2002-242022/29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21-AUG-2001; 2001WO-US026161
                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 129
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (SMIK ) SMITHKLINE BEECHAM CORP (SMIK ) SMITHKLINE BEECHAM PLC.
                                                                                                                                                                                                                                                                                                                              100;
                                                                                                                                                                                                                   64
                                                                                                                                                                                   64
                                                                                                                                                                                                                                                        4
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                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                     SGSGTSYSLTISRMEAEDAATYYCQQWSSNPLTFGSGTKLELKR 107
                                                                                                                                                                                                                                                      LTQSPAIMSASPGEKVTMTCSASSSVSYMYWYQQKPGSSPRLLIYDTSNLASGVPVRFSG
                                                                                                                                                                                                                                                                         LTQSBAIMSASPGEKVTTTCSASSSVSYMYMYQQXEGSSFRLLIYDTSNLASGVEVRFSG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Tornetta MA,
                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                  AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /note= "complementarity determining region 2"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /label= CDR3
/note= "complementarity determining region 3"
                                                                                                                                                                                                                                                                                                                                               93.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Truneh A,
                                                                                                                                                                                                                                                                                                                              2; Mismatches
                                                                                                                                                                                                                                                                                                                                               Score 516; DB 5;
Pred. No. 1.6e-31;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Wattam
                                                                                                                                                                                                                                                                                                                                                                5; Length 129;
                                                                                                                                                                                                                                                                                                                                2:
                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                  0;
                                                                                                                                                                                                                                                                                                                                Gaps
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RESULT 6
ADU86565
ID ADU8
XX
AC ADU8
XX
PT 10-F
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                                                                                                                                                                                                                                                                                                                                                                                                                          문
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CC call binding agent and a cytotoxic conjugate which comprises a CC call binding agent and a cytotoxic agent, where the call binding agent to CA6 glycotope. Preferably, the call binding agent is the murine CC anti-CA6 monoclonal antibody DS6 or their epitope-binding fragments or a CC humanized version of the same. The cytotoxic agent is chosen from CC maytansinoid compounds, taxoid compounds, CC-1065 compounds. The invention CC may be useful for the production of compounds with a cytostatic, CC immunosuppressive or virucide activity acting as stimulators of CC apoptosis. The invention is useful for inhibiting growth of a cell CC expressing CA6 glycotope and may be used for treating a subject having a CC cancer (for example serous ovarian carcinoma, endometriod ovarian CC carcinoma, neoplasm of the uterine cervix, neoplasm of the endometrius, neoplasm of the vulva, breast carcinoma, pancreatic tumor or tumor of the CC curchfelium), in which CA6 glycotope is expressed or overexpressed. In addition, the invention may be useful for treating autoimmune diseases, CC graft rejections or viral infections. The present sequence is that of a crejion of the mouse and protein which was used to demonstrate homology with a region of the mouse DS6 antibody light chain protein during the cx exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   apoptosis stimulation; cancer; ovary tumor; uterine cervix tumor; endometroid carcinoma; breast tumor; pancreas tumor; CA6 glycotope; autoimmune disease; graft rejection; viral infection; DS6;
             10-FEB-2005
                                                                           ADU86565;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 107 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 12; SEQ ID NO 46; 179pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cytotoxic agent.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cytotoxic conjugate useful for treating cancer, autoimmune diseases, graft rejections or viral infections, comprises cell binding agent and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2005-142544/15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Payne
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21-JUL-2004; 2004WO-US023340.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO2005009369-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            antibody engineering; ay1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cytotoxin; immunoconjugate; cytostatic; immunosuppressive; virucide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel
                                                                                                                                      ADU86565 standard; protein; 210 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (IMMU-) IMMUNOGEN INC
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                                                                                                                                                                                                                                                                                                    61
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        μ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 93.1%;
Similarity 91.6%;
                                                                                                                                                                                                                                                                                                                              FSGSGSGTSYSLTISRMEAEDAATYYCQQWSSNPLTFGSGTKLELKR 107
                                                                                                                                                                                                                                                                                                                                                                                                                          DTELTQSPAIMSASPGEKVTTTCSASSSVSYMYWYQQKPGSSPRLLIYDTSNLASGVPVR
                                                                                                                                                                                                                                                                                             FSGSGSGTSYSLTISRMEAEDAATYYCQQWSTYPLTFGAGTKLELKR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 515; DB 9;
Pred. No. 1.5e-31;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 107;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
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DTELTQSPAIMSASPGEKVTTTCSASSSVSYMYMYQQKPGSSPRLLIYDTSNLASGVPVR

DIOMTOSPAIMSASPGEKVTMTCSASSSVSYMYWYQQKPGSSPRLLIYDSTNLASGVPVR

8 60

FSGSGSGTSYSLTISRMBAEDAATYYCQQWSSNPLTFGSGTKLELKR 107

FSGSGSGTSYSLTISRMEAEDAATYYCQQWSTYPLTFGAGTKLELKR

107

믕 S

61 61

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CC reversibly from a repertoire of polypeptides that unfolds reversibly and comprises providing a polypeptides that unfolds from unfolded or misfolded bolypeptides. The method cc polypeptides from unfolded or misfolded bolypeptides. The method cc comprises providing a polypeptides; unfolding at least a portion of the unfolded cc polypeptides, and recovering at least one polypeptide that unfolds comprisibly and has the selectable characteristic from the refolded cc polypeptides, and recovering at least one polypeptide that unfolds creversibly and has the selectable characteristic from the refolded cc polypeptides, and recovering a polypeptide that unfolds creversibly from a repertoire of polypeptides that unfolds creversibly from a repertoire of polypeptides that unfolds reversibly. The clibrary or repertoire is useful for selecting a polypeptide comprising an antibody variable domain that unfolds reversibly or a polypeptide that cc refolds reversibly and comprising an antibody format. The polypeptide is useful in the manufacture of a medicament for diagnosing, treating or preventing a disease or medical condition mediated by a cytokine, cc cytokine receptor, enzyme, enzyme of action or DNA binding protein, such cc an inflammatory state, allergic hypersensitivity, cancer, bacterial or viral infection or an autoimmune disorder, e.g., asthma, psoriasis, Type c region associated with the protein purification method of the invention.
Query Match
Best Local S
Matches 98
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Recovering a polypeptide that unfolds reversibly from a repertoire polypeptides for treating e.g., cancer, by unfolding a portion of t displayed polypeptides and refolding a portion of the unfolded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     antidiabetic; vasotropic; vaccine; protein purification; protein foldin diagnosis; inflammation; immune disorder; allergic hypersensitivity; infection; autoimmune disease; asthma; psoriasis; insulin dependent disease; multiple sclerosis; rheumatoid arthritis; systemic lupus erythematosus; myasthenia gravis; hematological disease; systemic lupus erythematosus; myasthenia gravis; hematological disease;
                                                                                                                                             Sequence 210
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention describes a method of recovering a polypeptide that unfolds reversibly and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; SEQ ID NO 54; 222pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI, 2004-821888/81.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       14-MAY-2003; 2003US-0470340P
17-MAR-2004; 2004US-0554021P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           antiinflammatory; antibacterial; virucide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    polypeptides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO2004101790-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Immunoglobulin variable domain protein sėgid 54.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (DOMA-) DOMANTIS LTD
   186
                                     Similarity
   Conservative
                                                                                                                                                 8
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                                     93.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Famm
9
Score 515; DB
Pred. No. 2.9e
6; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          KHJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Winter GP;
                                     DB 8;
.9e-31;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cytostatic; antipsoriatic;
                                                                      Length 210;
       Indels
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   Gaps
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Best Loc
Matches
                   Query Match
 Local Similarity
         93.1%;
Score 515; DB 8; Length 210; Pred. No. 2.9e-31; 6; Mismatches 3; Indels
  0
  Gaps
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Sequence 108

Sequence

210 AA;

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RESULT 7
ADU8627
ADU8627
AX
AX
AX
AX
ADU86
AX
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                                                     CC comprises: providing a polypeptide display system comprising the cc repertoire of displayed polypeptides; unfolding at least a portion of the unfolded CC polypeptides; and recovering at least one polypeptide that unfolded CC polypeptides; and recovering at least one polypeptide that unfolded CC portion. The method is useful in recovering a polypeptide that unfolds CC reversibly and has the selectable characteristic from the refolded CC portion. The method is useful in recovering a polypeptide that unfolds comprising an antibody variable domain that unfolds reversibly. The CC useful in the manufacture of polypeptides that unfolds reversibly and comprising an antibody format. The polypeptide is CC useful in the manufacture of a medicament for diagnosing, treating or preventing a disease or medical condition mediated by a cytokine, CC useful infection or an autoimmune disorder, e.g., asthma, psoriasis, Type CC viral infection or an autoimmune disorder, e.g., asthma, psoriasis, Type CC region associated with the protein purification method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Recovering a polypeptide that unfolds reversibly from a repertoire of polypeptides for treating e.g., cancer, by unfolding a portion of the displayed polypeptides and refolding a portion of the unfolded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         14-MAY-2003; 2003US-0470340P.
17-MAR-2004; 2004US-0554021P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  systemic lupus erythematosus; myasthenia gravis; hematological disease; neoplasm; immunoglobulin; variable region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    antidiabetic; vasotropic; vaccine; protein purification; protein folding; diagnosis; inflammation; immune disorder; allergic hypersensitivity;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Immunoglobulin variable domain protein segid 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADU86571;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention describes a method of recovering a polypeptide that unfolds reversibly from a repertoire of polypeptides that unfolds reversibly and has a common selectable characteristic that distinguishes folded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2004-821888/81.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Jespers LS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         14-MAY-2004; 2004WO-GB002102.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     antiinflammatory; antibacterial; virucide; cytostatic; antipsoriatic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  polypeptides from unfolded or misfolded polypeptides. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; SEQ ID NO 60; 222pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (DOMA-) DOMANTIS LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Jones PC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
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ARESULT 8
ARESULT 8
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                                                                                This invention relates to the novel tredtment of an autoimmune disease chat utilises an antibody/ autoantigen donstruct. Specifically, it refers contains a peptide autoantigen that is linked to an antibody raised against an example of the presenting cell (APC) receptor. The present invention describes a contain antibody raised against an example of the relating diabetes mellitus by inducing an immune tolerance to an autoantigen, which can be chosen from glutamic acid decarboxylase (GAD), insulin or a heat shock protein (HSP), as well as epitopes derived from each thereof or beta cell antigens that are linked to an APC consent from DC-SIGNR, MHC, toll receptor or manose receptor amongst chers, which is expressed on the surfade of the APC. The antibody/ conters, which is expressed on the surfade of the APC. The antibody/ contents, which is expressed on the surfade of the APC. The antibody/ content then works to interfere with the interaction between the APC and immune cells such as autoreactive T cells, and hence contidiate to a proliferation. Accordingly, such compositions exhibit contidiates to interactions. Note that the single chain antibodies (scrye) used in this specification contain both variable light and heavy chain cregions connected by a linker, their short length makes these antibody contains to the receptor is preserved. This polypeptide sequence is a light chain murine anti-human L-SIGN (hL-SIGN) antibody protein of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            autoimmune disease;
diabetes mellitus; i
vaccine; hL-SIGN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 04-MAR-2003; 2003US-0451816P
15-DEC-2003; 2003US-0529500P
28-FEB-2004; 2004US-0548385P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Treating an autoimmune disease comprises providing an antibody/autoantigen construct containing an autoantigen linked to an antibody to a receptor of an antigen presenting cell and administering
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADT74993 standard; protein;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             04-MAR-2004; 2004WO-US006570
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 61
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   μ.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         9; Fig 8; 71pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Kretz-Rommel A,
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Length 108

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The invention describes an isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a specific clone where (I) binds to and neutralises botulinum neurotoxin type A (BONT/A). An isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a clone chosen from clone $25, C25, C39, IC6, 3D12, B4, IF3, huC25, Ar1, Ar2, WR1(V), WR1(T), 3-1, CC 3-8, 3-10 and ING1, Where (I) binds to and neutralizes botulinum composition (II) comprising BONT/A neutralising epitope having an epitope that is specifically bound by an antibody expressed by clones as mentioned in (I) is producing (I); and a composition (III) comprising several antibotulinum neurotoxin antibodies, where each antibody is specific for a contibodies shows greater toxin neutralisation than the single antibodies in surplus. The following are disclosed: a pharmaceutical composition (C comprising (I); and a kit comprising (I). (I) is useful for neutralising C showly and hit comprising (I). (I) is useful for neutralising C showly antibodies and a botulinum neurotoxin which (I) in surplus, where each of (I) is specific for a different epitope of the botulinum neurotoxin than the composition of involves contacting neurotoxin with (I) in surplus, where each of (I) is specific for a different epitope of the botulinum neurotoxin than the composition of antibodies shows greater toxin neutralisation than the composition of antibodies in surplus. (I) is useful for diagnosing the botulism correction than the composition of antibodies associated with botulinum neurotoxin than the composition than the composition of antibodies associated with botulinum neurotoxin than the composition than the composition of antibodies shows greater toxin neutralisation than the composition of antibodies shows greater toxin neutralisation than the composition than the composition than the composition of the composition of a composition of antibodies shows greater toxin neutralisation than the composi
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Best Local 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      useful for diagnosing with botulinum neurotoxin poisoning.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New isolated antibody that neutralizes botulinum neurotoxin type A, useful for diagnosing botulism or for treating pathologies associated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BONT/A neutralising epitope; anti-botulinum neurotoxin antib
toxin neutralisation; botulinum neurotoxin poisoning; mouse;
light chain variable region; single chain antibody; scFv.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2004-652009/63.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Marks JD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            31-AUG-1998; 98US-00144886.
01-AUG-2002; 2002US-0400721P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-AUG-2003; 2003US-00632706
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             09-SEP-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US2004175385-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              antibacterial; antibody; botulinum neurotoxin type A; BONT/A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mouse light chain variable region scrv seqid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       02-DEC-2004 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (REGC ) UNIV CALIFORNIA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LTQSPAIMSASPGEKVTMTCSASSSVSYMYWYQQKPGSSPRLLIYDTSNLASGVPVRFSG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Amersdorfer P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQ ID NO 72; 110pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 peptide;
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1; Mismatches
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Pred. No. 3.7e-31;
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RESULT 10
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Best Local S
Matches 98
                                                   Anti-epidermal growth factor receptor (EGFR) single chain antibodies and antibodies constructed from anti-EGFR antibody fragments can be used for diagnosis of tumours and assessment of tumour growth in vitro and in vivo. They may also be used in a pharmaceutical composition for the therapy of e.g. melanomas, gliomas and carcinomas. The antibodies and fragments are derived from mice but are humanised so as to cause minimum reaction against them. They are produced using the phage antibody library. They are produced using the phage antibody 1AAT04011-T04026 and AAR79858-R79873)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          poisoning. (I) exhibits specificity and affinity towards BoNT/A. enables rapid detection or diagnosis of botulism. This is the ami sequence of a mouse light chain variable fragment anti-botulinum scFv.
  Sequence 244
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   tumour; cancer; glioma; melanoma;
phage antibody library.
                                                                                                                                                                                                                                                                                                                                    Claim 4; Page 60-61; 93pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                    Anti-EGFR antibodies and single chain Fv antibody fragments - obtained from phage-antibody libraries, useful for diagnosis and therapy of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       N-PSDB; AAT04020
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Kettleborough AC,
Mitjans F, Rosell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               02-DEC-1994;
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                                                                                                                                                                                                                                                                                                                                                                                             tumours.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1995-336972/43.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (MERE ) MERCK PATENT GMBH
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l E, Blasco
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Pred. No. 7
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F, Piulats
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Query Match Best Local Similarity

91.5%; 92.6%;

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                                                                                                                                                                                                                             Query Match
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Matches 98
                                                                                                                                                                                                                                                                                                                                                                                             Anti-epidermal growth factor receptor (EGFR) single chain antibodies and antibodies constructed from anti-EGFR antibody fragments can be used for diagnosis of tumours and assessment of tumour growth in vitro and in vivo. They may also be used in a pharmaceutical composition for the therapy of e.g. melanomas, gliomas and carcinomas. The antibodies and fragments are derived from mice but are humanised so as to cause minimum reaction against them. They are produced using the phage antibody library. They are produced using the phage antibody library. (See
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Single chain antibody; antibody; epidermal growth factor receptor; EGPR; tumour; cancer; glioma; melanoma; carcinoma; diagnosis; assessment; phage antibody library.
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Mitjans F, Rosell E,
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                                                                                                                                                                                                                                                                                                                                               Sequence 242 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 4; Page 70; 93pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Anti-EGFR antibodies and single chain Fv antibody fragments - obtained from phage-antibody libraries, useful for diagnosis and therapy of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17-MAR-1994;
02-DEC-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FSGSGSGTSYSLTISRMEAEDAATYYCQQWSSYPPMYTFGGGTKLEIK 244
                                                                                                                                                DTELTQSDAIMSASPGEKVTTTCSASSSVSYMYWYQQKPGSSPRLLIYDTSNLASGVPVR 60
                           FSGSGSGTSYSLTISRMEAEDAATYYCQQWSSNPLTFGSGTKLELK 106
                                                                                                              DIELTQSPTIMSASPGEKVTMTCSDSSSVSYMYWYQQKTGSSPRLLIYDTSNLASGVPVR
FSGSGSGTSYSLTISRMEAEDAATYYCQQWSSYPLTFGAGTKLEIK 242
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PATENT GMBH.
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94EP-00118970.
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E, Blasco F,
                                                                                                                                                                                                                                                         91.3%;
                                                                                                                                                                                                                          Score 505; DB 2; Length 242;
Pred. No. 1.9e-30;
2; Mismatches 6; Indels
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F, Piulats J;
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LTQSPAIMSASPGEKVTTTCSASSSVSYMYWYQQKPGSSPRLLIYDTSNLASGVPVRFSG 63

Query Match
Best Local Similarity
Matches 97; Conserv

Conservative

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                                                           The invention discloses a modified antibody (or fragment) directed to the compidermal growth factor receptor (EGFR or Her 1). The antibody is communogenically non-immunogenic, or less immunogenic, than any original communogenically non-modified antibody directed to the same receptor when composed to the immune system of a given species and compared with the non condified antibody. Compared with a non-modified antibody, the modified compared with a non-modified antibody, the modified compared with a non-modified antibody. The same receptor when compared with a non-modified antibody. The same receptor when compared with a non-modified antibody. The same receptor when the mon-modified antibody. The consecutive amino acid residues of the 13-mer T-cell epitope having a consecutive amino acid residues of the 13-mer T-cell epitope having a consecutive amino acid residues of the 13-mer T-cell epitope having a consecutive amino acid residues of the 13-mer T-cell epitope having a consecutive amino acid residues of the 13-mer T-cell epitope having a consecutive amino acid residues of the 13-mer T-cell epitope having a consecutive aminopenically modified antibody having substantially no or less consecutive aminopenically modified antibody is useful in antibody therapy to treat cancers or consecutive modified antibody is useful in antibody therapy to treat cancers or consecutive modified antibody is substantially non-modified antibody of insecutive consecutive and with reduced or absent consecutive species, has enhanced properties and with reduced or absent consecutive and with reduced or absent consecu
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New modified anti-endodermal growth factor receptor antibodies which non-immunogenic or less immunogenic compared with non-modified antibodies, useful for treating cancers or tumors, e.g. tumor of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EGFR; antibody; Her 1; immunogenic; immune system; T cell epitop immunogenicity; major histocompatibility complex; MIC; allotype; MHC class II; antibody therapy; cancer; tumour; murine;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             breast, heart or lung.
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Hanlon M,
                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to an immunogenically modified fusion protein derived from a parent fusion protein, comprising first and second proteins/polypeptides, where the first protein is an immunoglobulin molecule or its fragment and the second protein is non-immunoglobulin target polypeptide each linked to the other directly or by a linker molecule. The immunogenically modified fusion protein is useful in combining known favourable properties of a composition or in creating new properties of a composition which elicits biological or pharmacological efficacy without having undesirable physiological effects such as nausea or gastric upset. The present sequence is mouse 425 antibody Vx protein. This sequence is used in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New modified fusion protein with reduced immunogenicity, useful combining favorable properties of a composition, comprises an immunoglobulin molecule linked to a non-immunoglobulin target
                                                                                                                                                                                                                                                                                                                                                  Sequence 106 AA;
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05-APR-2001; 2001EP-00108291.
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SGSGTSYSLTISRMEAEDAATYYCQQW8SHIFTFGSGTKLEIK 106
                                                     SGSGTSYSLTISRMEAEDAATYYCQQWSSNPLTFGSGTKLELK 106
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Watkins J,
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Baker M,
                                                                                                                                                                                                                           Score 502; DB 5; L
Pred. No. 1.5e-30;
Pred. No. 1.5e-30;
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RESULT 14
AAR27050
ID AAR277050
ID AAR277
XX AAR27
XX AAR27
XX 71425
XX VL425
XX VL425
XX WMONOC
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                                                                                                                                                                                                                                                                                                                                                           The variable light chain of monoclonal antibody 425 was prepd. from a synthetic cDNA sequence in which mutations had been made to the 5' and 3' ends to allow for cloning into HCMV expression vectors. Donor splice sites were recreated in the 3' flanking regions to allow correct splicing of the variable and constant regions. The 5' sequence was altered to introduce an initiation codon. The cloned antibody may be used int the prodn of reshaped or humanised antibodies which are less immunogenic than native antibodies and may be used to combat e.g. glioma, melanoma or carcinoma. See also AAR27299-300 and AARB7037-41. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human monoclonal antibodies binding to human receptors - and diagnosis of tumours, e.g. melanoma and carcinoma.
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N-PSDB; AAQ28740.
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01-MAR-1993
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                                                                                                    LTQSPAIMSASPGEKVTMTCSASSSVTYMYWYQQKPGSSPRLLIYDTSNLASGVPVRFSG
                                                                                                                             LTQSPAIMSASPGEKVTTTCSASSSVSYMYWYQQKPGSSPRLLIYDTSNLASGVPVRFSG
  SGSGTSYSLTISRMEAEDAATYYCQQWSSHIFTFGSGTKLEIK 128
                             SGSGTSYSLTISRMEAEDAATYYCQQWSSNPLTFGSGTKLELK 106
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tumour; melanoma; carcinoma; glioma; light; heavy; variable;
                                                                                                                                                                                                         Conservative
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71. .77
/label= CDR2
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13-FEB-1989;
28-SEP-1990;
19-DEC-1990;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention describes a method of producing humanised immunoglobulins involving expressing in a cell a nucleic acid encoding a humanised version of an immunoglobulin. This is obtained by comparing a donor and human immunoglobulin and producing a combined antibody which contains part of each. These are useful in the treatment of graft-versus-host disease, transplant rejection, autoimmune diseases such as diabetes, rheumatoid arthritis, myasthenia gravis, multiple sclerosis and systemic lugus erythematosus, herpes infections, (MV virus infections and myeloid leukaemia. The present sequence is an antibody used to demonstrate the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    07-JUN-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Producing humanized immunoglobulin, involves producing a cell containing DNA segments encoding humanized heavy and light chain variable regions, and expressing the DNA segments in the cell.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Fig 5; 145pp; English.
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89US-00310252.
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PL0013
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                               $03410
$66536
$26341
$29590
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A30562
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Description
g kappa chain V r
                                              A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-107 <SIKA
A;Cross-references: UNIPARC:UPI00001767B0
A;Cuperfamily: immunoglobulin V region; imm
C;Keywords: heterotetramer; immunoglobulin
F;16-89/Domain: immunoglobulin homology <IV
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A; Residues: 1-107 < DEN>
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   Query Match
Best Local Similarity
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    85.5%;
87.5%;
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438 79.2 132 2 S05268 Ig kappa chain pre 437.5 79.1 106 2 S11114 Ig kappa chain vr 437.7 79.0 130 2 S04573 Ig kappa chain vr 436 78.8 106 2 PS0070 Ig kappa chain vr 435 78.7 97 2 PH1084 Ig kappa chain vr 436 78.5 104 2 JC6076 Ig hight chain vr 437 78.3 108 2 PL0278 Ig kappa chain vr 438 78.3 108 2 PL0278 Ig kappa chain vr 439 78.1 130 2 B32456 Ig kappa chain pre 430 77.8 108 2 PS0069 Ig kappa chain vr 429 77.6 108 2 PL0277 Ig kappa chain vr 429 77.6 108 2 PL0276 Ig kappa chain vr 429 77.6 108 2 PL0276 Ig kappa chain vr 429 77.6 108 2 PL0276 Ig kappa chain vr 429 77.6 108 2 PL0276 Ig kappa chain vr 429 77.6 108 2 S11113 Ig kappa chain vr 420 77.2 107 2 S11113 Ig kappa chain vr 420 77.2 107 2 S11123 Ig kappa chain vr 421 77.2 107 2 S11123 Ig kappa chain vr 422 77.2 107 2 S11123 Ig kappa chain vr	u	4	ω	2	1	0	9	œ	7	0	ŋ	4.	ω	N		0
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2 S05268 Ig kappa 2 S11114 Ig kappa 2 S04573 Ig kappa 2 PS0070 Ig light 2 PH1084 Ig light 2 PH2078 Ig light 2 PL0278 Ig kappa 2 PL0278 Ig kappa 2 B32456 Ig kappa 2 B32456 Ig kappa 2 PL0277 Ig kappa 2 PL0277 Ig kappa 2 PL0277 Ig kappa 2 S11113 Ig kappa 2 S11113 Ig kappa 2 S11113 Ig kappa 2 S11113 Ig kappa	76.9	77.2	77.4	77.6	77.6	77.6	77.8	78.1	78.3	78.3	78.5	78.7	78.8	79.0	79.1	79.2
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Ig kappa chain pre Ig kappa chain V r Ig kappa chain V r Ig light chain V r Ig kappa chain V r	S05267	S11123	S11113	S29581	PL0276	PL0277	690084	B32456	S05269	PL0278	JC6076	PH1084	PS0070	S04573	S11114	S05268
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RESULT 2
A30562
Ig kappa chain V regions (27.7.2 and 27.4b.2) - mouse
Ig kappa chain V regions (27.7.2 and 27.4b.2) - mouse
C; Species: Mus musculus (house mouse)
C; Species: Mus musculus (house mouse)
C; Date: 23-Mar-1989 #sequence_revision 23-Mar-1989 #text_change 21-Jan-2000
C; Accession: A30562
C; Accession: A30562
R; Sikder, S.K.; Borden, P.; Gruezo, F.; Akolkar, P.N.; Bhattacharya, S.B.; Morrison, S.L.
R; Sikder, S.K.; Borden, P.; Gruezo, F.; Akolkar, P.N.; Bhattacharya, S.B.; Morrison, S.L.
R; Sikder, S.K.; Borden, P.; Gruezo, F.; Akolkar, P.N.; Bhattacharya, S.B.; Morrison, S.L.
R; Sikder, S.K.; Borden, P.; Gruezo, F.; Akolkar, P.N.; Bhattacharya, S.B.; Morrison, S.L.
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R; Sikder, S.K.; Borden, P.; Gruezo, F.; Akolkar, P.N.; Bhattacharya, S.B.; Morrison, S.L.
R; Sikder, S.K.; Borden, P.; Gruezo, F.; Akolkar, P.N.; Bhattacharya, S.B.; Morrison, S.L.
R; Sikder, S.K.; Bhattacharya, S.B.; Morrison, S.L.
R; Sikder, S.K.; Bhattacharya, S.R.; Bhattacharya, S.B.; Morrison, S.L.
R; Sikder, S.K.; Bhattacharya, S.R.; Bhattacharya, S.R.; Bhattacharya, S.R.; Bhattacharya, S.R.; Bhattacharya, S.R.; Bhattacharya, S.R.; Bhattach
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PC4405
Ig kappa chain V region (F3, anti-AFP) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 10-Nov-1997 #sequence_revision 23-Jan-1998 #text_(F)Cacession: PC4405
C;Accession: PC4405
C;Accession: PC4405
Chinese Biochem. J. 12, 648-653, 1996
A;Title: Generation of a phage display library of the imm
A;Reference number: PC4405
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A;Experimental source: spleen cell
C;Superfamily: immunoglobulin V region; immunoglobulin homology
F;16-89/Domain: immunoglobulin homology < NMM95
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PSGSGSGTSYSLTISRMEAEDAATYYCQQW9SNPLTFGSGTKLELKR
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Score 473; DB 2;
Pred. No. 4.4e-34;
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A;Cross-references: UNIPARC:UDI000017371F; GB:M27792; NID:g197159
A;Experimental source: mRNA clones for anti-phenyloxazolone antibody 6F6
A;Note: the authors translated the codon TrC for residue 8 as Pro and TrC for residue 8,Note: the codon given for 1-Met (AGT) is inconsistent with the authors' translation R;Milstein, C.; Even, J.; Jarvis, J.M.; Gonzalez-Fernandez, A.; Gherardi, E. Bur. J. Immunol. 22, 1627-1634, 1992
Bur. J. Immunol. 22, 1627-1634, 1992
A;Title: Non-random features of the repertoire expressed by the members of one V kappa A;Reference number: A49044; MUID:92289826; PMID:1601044
A;Accession: A49044
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ig kappa chain precursor V region (anti-phenyloxazolone 6F6) - mouse C;8pecies: Mus musculus (house mouse) C;Date: 31-Dec-1991 #sequence revision 09-Aug-1996 #text_change 16-Jul-1999 C;Accession: JL0079; A49044; B49044 R;Kaartinen, M.; Rocca-Serra, J.; Maekelae, O. Mol. Immunol. 25, 859-865, 1988 A;Title: Combinatorial association of V genes: one VH gene codes for three rapression: JL0076; MUID:89096973; PMID:3211160
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A;Cross-references: UNIPARC:upI00001767BA
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;16-89/Domain: immunoglobulin homology <IMM>
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C;Species: Mus musculus (house mouse)
C;Date: 23-Mar-1989 #sequence_revision 23-Mar-1989 #text_change 21-Jan-2000
C;Accession: B30562
R;Sikder, S.K.; Borden, P.; Gruezo, F.; Akolkar, P.N.; Bhattacharya, S.B.; I
J. Immunol. 142, 888-893, 1999
                                                                  A; NOLECULE type: DNA
A; Molecule type: DNA
A; Residues: 1-25 <MIL>
A; Residues: 1-25 <MIL>
A; Cross-references: UNIPARC: UPI000011760C; GB:S37663; NID:g250214; PIDN:AAB22331.1;
A; Cross-references: UNIPARC: UPI000011760C; GB:S37663; NCBIN:106802, NCBIP:106809)
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                          A; Molecule type: DNA
A; Residues: 114-116 < MIL2>
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A;Status: preliminary
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A; Residues: 1-130 < KAA>
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;Cross-references: UNIPARC:UPI000011E8E5; GB:S37664; NID:g250215; PIDN:AAB22332.1;
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Best Local 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SGSGTSYSLTISRMEAEDAATYYCQQWSSNPLTFGSGTKLELKR 107
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86.5%;
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Pred. No. 5.3e-34;
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Ig light chain V region - mouse C; Species: Mus musculus (house mouse) C; Date: 06-Jan-1995 #sequence_revision 06 C; Accession: S38720 R; Cimanis, A.Y. submitted to the EMBL Data Library, NovembarReference number: S38713
                                                                                                                                               Ig kappa chain V region - mouse
C;Species: Mus musculus (house mouse)
C;Date: 06-Jan-1995 #sequence_revision
C;Accession: 829591
R;Kaveler, J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 5
S38720
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A;Note: sequences extracted from NCBI backbone; partial nucleotide sequences of 1
A;Note: sequence extracted from NCBI backbone (NCBIN:106807, NCBIP:106822)
C;Genetics:
A;Gene: V(kappa)Ox1
A;Introns: 17/1
C;Complex: An immunoglobulin heterotetramer subunit consists of two identical light of complex: An immunoglobulin heterotetramer subunit consists of two identical light of cyling in the subunit sessociate c;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;13-130/Product: Ig kappa chain V region (6F6) #status predicted <MAT>
F;38-111/Domain: immunoglobulin homology
                                                            R;Kavaler, J.
submitted to the EMBL Data
A;Reference number: 826459
A;Accession: 829591
A;Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Status: preliminary
A;Molecule type: mRNA
A;Molecule type: mRNA
A;Residues: 1-108 <CIM>
A;Residues: 1-108 <CIM>
A;Cross-references: UNIPARC:UPI0000117544; EMBL:X76023; NID:g416104;
A;Cross-references: UNIPARC:UPI0000117544; EMBL:X76023; NID:g416104;
C;Superfamily: immunoglobulin V region; immunoglobulin homology <IMM>
F;16-89/Domain: immunoglobulin homology <IMM>
A;Molecule type: mRNA
A;Residues: 1-103 <KAV>
A;Cross-references: UNIPARC:UPI0000115F57; EMBL:X59094; NID:g52227; PIDN:CAA41820.1; PID:
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S29591
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           93,
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Similarity 88.6%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                     LTQSPALMSASPGEKVTMTCSASSSVSYMYWYQQKPR88PKPWIYLTSNLASGVPLRF8G
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 469.5; DB 2; Pred. No. 8.8e-34;
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Pred. No. 7.9e-34;
                                                                                                                             April
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A;Status: preliminary
A;Molecule type: mRNA
A;Molecule type: mRNA
A;Residuse: 1-235 <FIS>
A;Residuse: 1-235 <FIS>
A;Residuse: 1-205 <FIS>
A;Cross-references: UNIPARC:UPI0000116096; EMBL:X67211; NID:g54828; PIDN:CAA47650.1; PID
C;Superfamily: immunoglobulin V region; immunoglobulin homology
F;38-111/Domain: immunoglobulin homology <IMM>
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A;Accession: $25058
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C; Keywords: het
F; 15-88/Domain:
                                                                                                                                                                                                                                                                                                    R;Levy, S.; Campbell, M.J.; Levy, R.
J. Exp. Med. 170, 1-13, 1989
A;Title: Functional immunoglobulin light chain genes are replaced A;Reference number: A92781; MUID:89310348; PMID:2501443
A;Accession: PS0071
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          R;Fischer, R.; Voss, A.; Niersbach, M.; Munziker, W.; Hirsch, H.J.; Kreuzaler, submitted to the EMBL Data Library, July 1992
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A; Residues: 1-106 < LEV>
                                                                                                                                                                                                                                                                                  A; Status: translation not shown
                                                                                                                                                                                                                                                                                                                                                                                                                                   Ig kappa chain V region (38C13.V8) - mouse
C;Species: Mus musculus (house mouse)
C;Date: 31-Mar_1990 #sequence_revision 31-Mar-1990 #text_change 21-Jan-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Species: Mus musculus (house mouse)
;Date: 19-Mar-1997 #sequence_revision 19-Mar-1997 #text_change 21-Jan-2000
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Best Local :
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Best Local
                                                                                                                                                     ;Cross-references: UNIPARC:UPI00001767C9

;Superfamily: immunoglobulin V region; imm

;Keywords: heterotetramer; immunoglobulin

;16-89/Domain: immunoglobulin homology <IN
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LTQSPAIMSASPGEKVTTTCSASSSVSYMYWYQQKPGSSPRLLIYDTSNLASGVPVRFSG
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Pred. No. 3.9e-33;
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Pred. No. 1.4e-33;
                                                                                      Score 465; DB 2;
Pred. No. 2.1e-33;
                                                                      Mismatches
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Eur. J. Immunol. 22, 2893-2899, 1992
A;Title: Molecular analysis of anti-idiotypic monoclonal antibodies in the HLA-DR antiger
A;Reference number: A49049; MUID:93049629; PMID:1425914
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C;Species: Mus musculus (house mouse)
C;Date: 19-Dec:1993 #sequence_revision 18-Nov-1994 #text_change 21-Jan-2000
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A;Experimental source; cell line 4Cl1
C;Comment: This protein is an anti-idiotypic;antibody that induces an anti-phosphorylchol C;Comment: This protein is an anti-idiotypic;antibody that induces an anti-phosphorylchol C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
C;Keywords: heterotetramer; immunoglobulin
C;Keywords: heterotetramer; immunoglobulin
C;Keywords: heterotetramer; immunoglobulin V region (4G11) #status predicted <MAT>
F;1-22/Domain: immunoglobulin homology <INM>
F;33-112/Domain: immunoglobulin homology <INM>
F;36-55/Region: complementarity-determining 1
F;71-77/Region: complementarity-determining 2
F;71-77/Region: complementarity-determining 1
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A;Title: Structural basis of stimulatory anti-idiotypic antibodies.
A;Reference number: PL0011; MUID:88142863; PMID:3125424
A;Accession: PL0013
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ig kappa chain precursor V region (4C11) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 30-Jun-1992 #sequence_revision 30-Jun-1992 #text_change 21-Jan-2000
C;Accession: PL0013
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PL0013
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A; Residues: 1-104 < ARM>
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R;Armandola, E.A.; Mariani, S.M.; Zwickl, M.; Hardman, N.; Ferrone,
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A; Residues: 1-140 < CHE>
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Best Local
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                                                                                                                  Best Local
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Similarity 88.1%;
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LTQSPAIMSASPGEKVTTTCSASSSVSYMYWYQQKPGSSPRLLIYDTSNLASGVPVRFSG
                                                                                                                                                                                                   complementarity-determining 3 constant region (fragment) #status predicted <COR>
                                                                                                                  87.5%;
                                                                                 Score 462; DB 2;
Pred. No. 5.1e-33;
5; Mismatches 8
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Pred. No. 3.1e-33;
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A;Molecule type: DNA
A;Residues: 1-107 <FEL>
A;Residues: 1.107 <FEL>
A;Cross-references: UNIPARC:UPI0000115334; EMBL:M90690
A;Cross-requence extracted from NCBI backbone (NCBIN:109958, NCBIP:109959)
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: immunoglobulin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         N;Alternate names: L6 anti-tumor antibody
C;Species: Mus musculus (house mouse)
C;Species: Mus musculus (house mouse)
C;Date: 27-Apr-1993 #sequence revision 18-Nov-1994 #text_change 21-Jan-2000
C;Accession: A42848; S33902
R;Fell, H.P.; Gayle, M.A.; Yelton, D.; Lipsich, L.; Schieven, G.L.; Marken,
J. Biol. Chem. 267, 15552-15558, 1992
A;Title: Chimeric L6 anti-tumor antibody. Genomic construction, expression,
A;Reference number: A42848; MUID:92348410; PMID:1639794
A;Accession: A42848
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C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
C;Keywords: heterotetramer; immunoglobulin
R;16-89/Domain: immunoglobulin homology <IMM>
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J. Immunol. 142, 863-870, 1989
A;Title: Variable region cDNA sequences and antigen binding
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C;Species: Mus musculus (house mouse)
C;Date: 23-Mar-1989 #sequence_revision 23-Mar-1989 #text_change
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A; Residues: 1-108 < MAT>
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Best Local S
Matches 89
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                                        SGSGTSYSLAISRVEABDAATYYCQQWNSNPLTFGAGTKLELKR
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                                                                                                                            LSQSPAILSASPGEKVTLTCRASSSVSFMNWYQQKPGSSPKPWIYATSNLASGVPGRFSG
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85.6%;
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                                                                                                                                                                                                                                  Score 460; DB 2;
Pred. No. 5.7e-33;
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Pred. No. 5.2e-33;
5; Mismatches
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Chinese BioChem. J. 12, 648-653, 1996
A; Title: Generation of a phage display library o
A; Reference number: PC4405
A; Contents: Spleen
A; Accession: PD0011
A; Molecule type: mRNA
A; Residues: 1-107 < DEN>
A; Cross-references: UNIPARC: UPI00001767AA
C; Superfamily: immunoglobulin V region; immunogl
F; 16-89/Domain: immunoglobulin homology < IMM>
                                 BSSULT 15
B54378
Ig light chain V region anti-triplex DNA - mouse (fragment)
Ig light chain V region anti-triplex DNA - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Species: Mus musculus (house mouse)
C;Date: 06-Oct-1994 #sequence_revision 18-Nov-1994 #text_change 21-Jan-2000
C;Date: 06-Oct-1994 #sequence_revision 18-Nov-1994 #text_change 21-Jan-2000
C;Accession: B54378
R;Agazie, Y.M.; Lee, J.S.; Burkholder, G.D.
J. Biol. Chem. 269, 7019-7023, 1994
J. Biol. Chem. 269, 7019-7023, 1994
A;Title: Characterization of a new monoclonal antibody to triplex DNA and immunofluoresce
A;Reference number: A54378; MUID:94165109; PMID:7509814
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Cross-references: UNIPARC:UPI00001767E0
C;Superfamily: immunoglobulin V region; immun
F;16-89/Domain: immunoglobulin homology <IMM>
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S11119
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A; Residues: 1-107 < NAT>
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Best Local S
Matches 88
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Best Local Similarity
Matches 88; Conserv
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Ig kappa chain V region (VLB10, anti-AFP) - mbuse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 10-Jul-1998 #sequence_revision 10-Jul-1998 #text_change 21-Jan-2000
C;Accession: PD0011
                                                                                                                                                                                                                                                                                                                                                                                                         A;Title: mRNA sequences define an unusually restricted IgG response to 2-phenyloxazolone A;Reference number: S07331; MUID:83271467; PMID:6877353 A;Accession: S11119
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                         SGSGTSYSLTISRMEAEDAATYYCQQWSSNPLTFGSGTKLELKR 107
                                                                                                                   LTQSPAIMSASPGEKVTTTCSASSSVSYMYWYQQKPGSSPRLLIYDTSNLASGVPVRFSG
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XGXGTSYSLTISSMFAEDAATYYCQQWSSNPLTFGAGTKLELKR
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A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-106 <AGA>
A;Cross-references: UNIFARC:UPIO0001767AC; GB:S68985; NID:g545746; PIDN:AAB30096.1; PID:
A;Cross-references: UNIFARC:UPIO0001767AC; GB:S68985; NID:g545744175)
C;Styperfamily: immunoglobulin v region; immunoglobulin homology
C;Keywords: immunoglobulin
P;16-89/Domain: immunoglobulin homology <IMM>
C;Keywords: immunoglobulin
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4 LTQSPAIMSASPGEKVTTTCSASSSVSYMYMYQQKPGSSPRLLIYDTSNLASGVPVRPSG 	puery Match 90.4%; Score 500; DB 2; Length 106; sest Local Similarity 94.2%; Pred. No. 3.6e-42; fatches 97; Conservative 2; Mismatches 4; Indels	SMR; Q9U410; 4-106. InterPro; IPR007110; Ig-like. InterPro; IPR007516; Ig_v. SMART; SM00406; IGv; 1. PROSITE; PS50835; IG_LIKE; 1. NON_TER 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	nd sequence analysis of the light chain variable oclonal anti-idiotypic antibody NP30 of Schistoso", Sheng Chong Xue Yu Ji Sheng Chong Bing Za Zhi 2000).  ANF19434.1; -; Genomic_DNA.	DUENCE.  DUENCE.  27;  19 Z.Q., Qiu Z.N., Li Y.Q., Yu X.C., Xiong Y.,  Jan X.H.;	Mus musculus (Mouse). Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleos Mammalla; Eutheria; Euarchontoglires; Glires; Rodentia; Sciu Muridae; Murinae; Mus.	2000 (TrEMBLrel. 13, Created) -2000 (TrEMBLrel. 13, Last sequence update) -2003 (TrEMBLrel. 25, Last annotation update) nal anti-idiotypic Schistosoma japonicum antibody globulin light chain variable region (Fragment).	PRELIMINARY;	ALIGNMENTS	354 64.0 107 2 353 63.8 109 1 352.5 63.7 108 2 352.5 63.7 111 1	9 354.5 64.1 111 1 KV3O MOUSE 0 354.5 64.1 236 2 Q6FILB HUMAN 1 354.5 64.1 240 2 Q65ZC9_HUMAN	7 354.5 64.1 108 2 Q9UL <sup>7</sup> 0_HUMAN	4 357.5 64.6 108 1 KV1B HŪMAN 5 355.5 64.3 108 1 KV1O HŪMAN 5 355.5 64.3 101 1 KV1M MOTER	2 358.5 64.8 111 1 KV3N MOUSE P0166
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01-OCT-2002 (TrEMBLrel. 26, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Anti-VIPase light chain variable region (Fragment).
Mus musculus (Mouse).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalla, Eutheria, Euarchontoglires, Glires, Rodentia, Sciurognal
Muridae, Muilnae, Mus.
    NUCLBOTIDE SEQUENCE.
STRAIN-BALB/C; TISSUE-Hyperimmunized spleen;
Zhou Y.-X., Taguchi H., Planque S., Karle S.,
Submitted (MAY-2002) to the EMBL/GenBank/DDBJ
[2]
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This Swiss-Prot entry is copyright. It is produced through a c
between the Swiss Institute of Bioinformatics and the EMBL
the European Bioinformatics Institute. There are no restrict
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Mus musculus (Mouse).
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                                                                                                                                                                                                                                                                                                                                             QBK1F2_MOUSE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SMR; P04945; 1-108.
Ensembl; ENSMUSG00000062686; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             removed
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Muroidea; Muridae; Murinae; Mus.
                                                                                                                              NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PROSITE, PS50835;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUCLEOTIDE SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        SGSATSYSLTITRMQAEDAATYYCQQWSSYPPMLTFGAGTKLELK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LTQSPAINSASPGQKVTMTCSASSSVSYMYWYQQKPGSSPRLLIYDTSNLASGVPVRFSG
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Euarchontoglires; Glires;
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Pred. No. 4.7e-41;
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Framework-3.
Complementarity-determining-3.
Framework-4.
By similarity.
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Complementarity-determining-1.
Framework-2.
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lires; Rodentia; Sciurognathi;
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                                                                                                                                                                        Sciurognathi;
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RESULT TO SEE AND SEE 
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                                                                                                                                                     RX MEDLINE=22386257; PubMed=12477932; DOI=10.1073/pnas.242603899;
RX Strausberg R.L., Feingold E.A., Grouse L., H., Derge J.G.,
RX Alausher R.D., Collins F.S., Wagner L., Bhenmen C.M., Schuler G.D.,
RX Altschul S.F., Zeeberg B., Buetow K.H., Bchaefer C.F., Bhat N.K.,
RX Altschul S.F., Zeeberg B., Buetow K.H., Bchaefer C.F., Bhat N.K.,
RX Altschul S.F., Zeeberg B., Buetow K.H., Bchaefer C.F., Bhat N.K.,
RX Altschul S.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RX Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RX Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RX Altschako L., Marusina K., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RX Brownstein M.J., Ugdin T.B., Toshiyuki S., Carninci P., Prange C.,
RX Brownstein M.J., Ugdin T.B., Toshiyuki S., Carninci P., Prange C.,
RX Brownstein M.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
RX Asha S.S., Loquellano N.A., Peters G.J., Abrameon R.D., Mullahy S.J.,
RX Raha S.S., Loquellano N.A., Peters G.J., Abrameon R.D., Hulay S.W.,
RX Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RX Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RX Richards S., Worley K.C., Hale S., Garcia A., Rodrigues S., Sanchez A.,
RX Willalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RX Wilting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RX Wilting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RX Milting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RX Butterfield Y.S.N., Truchman J.W., Green E.D., Dickson M.C.,
RX Schein J.E., Jones S.J.M., Marar M.A.,
RX Schein J.E., Jones S.J.M., Marar M.A.,
RX Schein J.E., Jones S.J.M., Marar M.A.,
RX Generation and initial analysis of more than 15,000 full-length human
RT and mouse cDNA sequences ";
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Best Local S
Matches 93
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10-MAY-2005 (TrEMBLrel. 30,
10-MAY-2005 (TrEMBLrel. 30,
10-MAY-2005 (TrEMBLrel. 30,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PROSITE;
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InterPro; IPR007710; Ig-like.
InterPro; IPR003596; Ig-v.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 1.
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Proc. Natl. Acad. Sci. U.S.A. 86:4624-4628(1989)
EMBL; AF516283; AAM64201.1; -; Genomic_DNA.
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STRAIN=FVB/N; TISSUE
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Mammalia; Eutheria; Euarchontoglires;
Muroidea; Muridae; Murinae; Mus.
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HSSP; P01837; 25C8.
SMR; Q8K1F2; 2-112.
NUCLEOTIDE SEQUENCE.
STRAIN=FVB/N; TISSUE=Colon;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Baccala R., Quang T.V., Gilbert M., Ternynck T., Avrameas S "Two murine natural polyreactive autoantibodies are encoded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mus musculus (Mouse)
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112 AA, 11953 MW,
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Last annotation update)
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Submitted (MAR-2005) to the EMBL; BC091738; AAH91738.1; SMR; Q58EV6; 23-235.

-; mRNA.

EMBL/GenBank/DDBJ

databases

Project;

GO; GO:0003823; F:antigen binding; IEA.

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Best Local S
Matches 90
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Interpro; IPR003599; Ig.
Interpro; IPR003597; Ig.Cl.
Interpro; IPR003597; Ig.Cl.
Interpro; IPR003596; Ig.MC.
Interpro; IPR003596; Ig.V.
Pfam; PP07654; Cl-set; 1.
SMART; SM00409; IG: 2.
SMART; SM00409; IG: 1.
SMART; SM00406; IGV; 1.
PROSITE; PS50835; IG LIKE; 2.
PROSITE; PS50835; IG MHC; UNKNOWN 1.
SEQUENCE 235 AA; 25719 MW; BE4B4AAB
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P04940;
13-AUG-1987 (Rel. (
13-AUG-1987 (Rel. (
10-MAY-2005 (Rel. (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUCLEOTIDE SEQUENCE.

MEDLINE=83271467; PubMed=6877353;

Kaartinen M., Griffiths G.M., Markham A.F., Milstein C.

"mRNA sequences define an unusually restricted IgG resp
phenyloxazolone and its early diversification.";

Nature 304:320-324(1983).

-i- FUNCTION: Anti-2-phenyl oxazolone (PHOX) Antibody.
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rkusiTB; PS50835; IG LIKB; 1.

Hybridoma; Immunoglobulin domain; Immunoglobulin REGION 1 23 Complementarity-dete. REGION 24 33 Complementarity-dete. REGION 34 48 Framework-2.

REGION 49 55 Complementarity-detes. Complementarity-detes. REGION 56 87 Framework-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOUSE
                                                                                                                                                                                                                                                                                                                                                                                                                              This Swiss-Prot entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on it use as long as its content is in no way modified and this statement is no
                                                                                                                                                                      Ensembl; ENSMUSG00000062047;
InterPro; IPR0077110; Ig-like.
InterPro; IPR003596; Ig-v.
SMART; SM00406; IG-v; 1.
PROSITE; PS50835; IG-LIKE; 1.
                                                                                                                                                                                                                                                                                                   EMBL; K00735; AAA38680.1;
HSSP; P01679; 2FBJ.
SMR; P04940; 1-107.
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el. 05, Last sequence update)
el. 47, Last annotation update)
V-VI region NQ2-17.4.1.
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Pred. No. 7.6e-39;
                                                                                                                                                                                                                                                                                Mus musculus.
                   Framework-2.
Complementarity-determining-2
Framework-3.
                                                                                               Complementarity-determining-1.
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P3 MOUSE
Q8K1F3
Q8K1F3
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PIR; PH1085; PH1085.

HSSP; P01837; 25C8.

SMX; Q8K1F3; 2-112.

Ensembl; ENSWISG00000063156; Mus m.

InterPro; IPR007110; Ig-like.

InterPro; IPR007596; Ig-v.

SMART; SM00406; IGV; 1.
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01-OCT-2002 (TrEMBLrel. 22, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Anti-VIPase light chain variable region (Fragment).
Mus musculus (Mouse).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognat
Muridae; Mus.
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SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                   MEDLINE=92381444; PubMed=1512540; DOI=10.1084/jem.176.3.761; Tillman D.M., Jou N.T., Hill R.J., Marion T.N.; Tillman D.M. Jou N.T., Hill R.J., Marion T.N.; Both IgM and IgG anti-DNA antibodies are the products of clesselective B cell stimulation in (NZB x NZW)FI mice."; J. Exp. Med. 176:761-779(1992).

TEMBL, AF516282; AAM64200.1; -; Genomic_DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Baccala R., Quang T.V., Gilbert M., Ternynck T., Baccala R., Quang T.V., Gilbert M., Ternynck T., "Two murine natural polyreactive autoantibodies nonmutated germ-line genes."; proc. Natl. Acad. Sci. U.S.A. 86:4624-4028(1989)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STRAIN=BALB/c; TISSUE=Hyperimmunized spleen; Zhou Y.-X., Taguchi H., Planque S., Karle S. Submitted (MAY-2002) to the EMBL/GenBank/DDB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUCLEOTIDE SEQUENCE
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                                                                                                                                                                                                         PROSITE;
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             1-4
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112 AA,
                                                      83.7%;
nilarity 85.0%;
Conservative '
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ne EMBL/GenBank/DDBJ
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                                                        Score 463; DB
Pred. No. 1.9e
4; Mismatches
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pred. No. 9.2e-39;
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                                                                                                                                              22DD75B5414CDB18 CRC64;
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autoantibodies ;
                                                                                                                                                                                                                                                                                                  musculus
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13-AUG-1987
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MEDLINE=83271467, PubMed=6877353;

MEDLINE=83271467, MISSELECT C.

MEDLINE=83271467, PubMed=6877353;

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13-AUG-1987 (Rel. 05, Created)
13-AUG-1987 (Rel. 05, Last sequence up
10-MAY-2005 (Rel. 47, Last annotation
19 kappa chain V-VI region NQ6-8.3.1.
Mouse)
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InterPro; IPR003596; Ig_v.
SMART; SM00406; IGv; 1.
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Ensembl; ENSMUSG00000062047; Mus
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H8SP; P01679; 2FBJ.
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Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;
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1 23 Framework-1.
24 33 Complementarity-determining-1.
34 48 Framework-2.
49 55 Complementarity-determining-2.
56 87 Framework-3.
88 96 Complementarity-determining-3.
87 Pramework-1.
97 106 Framework-1.
97 107 107
107 107
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Complementarity-determining-3.
Framework-4.
By similarity.
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Complementarity-determining-1.
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SOLUTION SERVICE COUNTY AND SERVICE SE
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I Ig kappa chain V-VI region NQ5-61.1.2.
Mus musculus (Mouse).
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Mammalia; Eutheria; Euarchontoglires; Gli
Murcidea; Murdae; Murinae; Mur
NUCLEOTIDE SEQUENCE.
MEDILINE=83271467; PubMed=6877353;
MEDILINE=83271467; PubMed=6877353;
Kaartinen M., Griffithe G.M., Markham A.F., Mils
"mRNA sequences define an unusually restricted I
phenyloxazolone and its early diversification.";
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HSSP; Q91W12; 1AY1.
SMR; P04941; 1-103.
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Mammalia; Eutheria; Buarchontoglires; Glires; Rodentia; Sciurognathi;
Muroidea; Muridae; Musiae; Mus.
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13-AUG-1987 (Rel. 05, Last
10-MAY-2005 (Rel. 47, Last
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InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
SMAXT; SM00406; IGv; 1.
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13-AUG-1987
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Mus musculus (Mouse)
                                                                                                                                                              NCBI_TaxID=10090;
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    -i- FUNCTION: Anti-2-phenyl oxazolone (PHOX) Antibody.

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Framework-4.
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InterPro; IPR003596; Ig v.
SMART; SM00406; IGv; 1.
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SMR; P04942; 1-107
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                                                                                                                                                                                                                             MUCLEOTIDE SEQUENCE.
MEDLINE-83271467; PubMed-6877353;
KABRITINE-83271467; PubMed-6877353;
KABRITINE M., Griffiths G.M., Markham A.F., Milstein
"MRNA sequences define an unusually restricted IgG re
phenyloxazolone and its early diversification.";
                                                                                                                                                                                                                                                                                                                                                                           Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Butheria, Euarchontoglires, Glires, Rodentia, Sciurognathi,
Muraldea, Muridae, Murinae, Mus.
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13-AUG-1987 (Rel.
10-MAY-2005 (Rel.
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EMBL; K00744; AAA38689.1;
HSSP; Q91W12; 1AY1.
SMR; P04944; 1-106.
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                                                                                                                                                                                          Nature 304:320-324(1983).
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FUNCTION: Anti-2-phenyl oxazolone (PHOX) Antibody.
                                                                                                                                   Swiss-Prot entry is copyright. It is produced through een the Swiss Institute of Bioinformatics and the EV
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Framework-3.
Complementarity-determining-3.
Framework-4.
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Pred. No. 3.7e-38;
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RA Altechul S.F., Zeebberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Altechul S.F., Zeebberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Altechul S.F., Zeebberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Hopkins R.F., Jordan H., Moore T., Max 91., Wang J., Hsieh F.,
RA Biatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casaninci P., Prange C.,
RA Brownstein M.J., Uddin T.B., Toshiyuki 9., Carninci P., Prange C.,
RA Brownstein M.J., Uddin T.B., Toshiyuki 9., Carninci P., Prange C.,
RA Brownstein M.J., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gunaratne P.H.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gunaratne P.H.,
RA Richards J., Helton B., Ketteman M., Madan A., Rodrigues S., Sanchez A.,
RA Pahey J., Helton B., Ketteman M., Madan A., Rodrigues S., Sanchez A.,
RA Willalon D.K., Mazny D.M., Sodergren B.J., Lu X., Gibbs R.A.,
RA Wilting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Mitting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Mitting M., Wadan A., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RGeneration and initial analysis of mord than 15,000 full-length human
"Generation and initial analysis of mord than 15,000 full-length human
and mouse cDNA sequences.",
RT proc. Natl. Acad. Sci. U.S.A. 99:16899-16903 (2002).
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Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia;
Muroldea; Muridae; Murinae; Muß.
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InterPro; iPR003596; ig_v.
SMART; SM00406; iGv; 1.
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25-OCT-2004
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Q5XFY8;
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STRAIN-FVB/N; TISSUE-Colon;
Director MGC Project;
Submitted (OCT-2004) to the
EMBL; BC084683; AAH84683.1;
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                                                             NUCLEOTIDE SEQUENCE
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Immunoglobulin c
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Last sequence update)
Last annotation update)
                EMBL/GenBank/DDBJ databases
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Pred. No. 3.7e-38;
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GO:0003823; F:antigen binding;

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RESULT
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Best Local S
Matches 87
                                                                         Query Match
Best Local 9
                                                      Matches
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InterPro; IPR007110; Ig-like.
InterPro; IPR003597; Ig_cl.
InterPro; IPR003597; Ig_MHC.
InterPro; IPR003006; Ig_MHC.
InterPro; IPR003566; Ig_v.
Pfam; PF07654; Cl.=set; l.
SMART; SM00409; IG; 2.
SMART; SM00407; IGcl; 1.
SMART; SM00407; IGcl; 1.
SMART; SM00407; IGcl; 1.
SMART; SM00408; IGv; I.
PROSITE; PS00290; IG_MHC; UNKNOWN_1.
                                                                                                                                                                                                                                                                                                                                                               PIR; A33933; A33933.
PIR; PC4405; PC4405.
HSSP; P01837; 10RQ.
SMR; Q8K1F0; 3-112.
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OBKIFO;

OBKIFO;

O1-OCT-2002 (TrEMBLrel. 22, Created)

O1-OCT-2002 (TrEMBLrel. 22, Last sequence update)

O1-MAR-2004 (TrEMBLrel. 26, Last annotation update)

O1-MAR-2004 (TrEMBLrel. 26, Last annotation update)

Anti-VIPasse light chain variable region (Fragment).

Mus musculus (Mouse).

Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Mammalia; Butheria; Euarchontoglires; Glires; Rodentia; Sciurognat

Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        "Two murine macutary Fractions of the genes.";
nonmutated germ-line genes.";
Proc. Natl. Acad. Sci. U.S. A. 86:4624-4628(1989).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUCLEOTIDE SEQUENCE.

STRAIN-BALB/c; TISSUE-Hyperimmunized spleen;
STROU Y.-x., Taguchi H., Planque S., Karle S.,
Submitted (MAY-2002) to the EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hypothetical protein. SEQUENCE 235 AA; 2
                                                                                                                                                                               PROSITE; PS50835; IG_LIKE; 1.
NON_TER 1 1 1
NON_TER 112 112
                                                                                                                                                                                                                                                      Ensembl; ENSMUSG00000062047; Mus musculus. InterPro; IPR007110; Ig-like. InterPro; IPR003596; Ig_v. SMART; SM00406; IGv; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Baccala R., Quang T.V., Gilbert M., Ternynck T., "Two murine natural polyreactive autoantibodies
                                                                                                                                                          SEQÜENCE
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                                                                           Similarity
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LTQSPAIMSASPGEKVTTTCSASSSVSYMYWYQQKPGSSPRLLIYDTSNLASGVPVRFSG
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112 AA;
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                                                                                                                                                       11901 MW; F6644663201AA239 CRC64;
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83.7%;
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                                                                           82.5%;
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                                                   Score 456; DB 2; 1
Pred. No. 9.8e-38;
4; Mismatches 12;
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Pred. No. 1.9e-37;
B; Mismatches 9
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RESULT 14
Q811C3_MOUSE P
ID Q811C3_MOUSE P
AC Q811C3_F
DT 01-UUN-2003 (T
DT 01-MAR-2004 (T
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Best Local (
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Ensembl; ENSMUSG0000062047; M.

MGI; MGI: 2686348; Gm1502.

InterPro; IPR0077110; Ig-1ike.

InterPro; IPR003596; Ig_v.

SMART; SM00406; IGv; 1.

PROSITE; P850835; IG LIKE; 1.

NON_TER

134

134
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01-MAR-2002 (TrEMBLrel. 20, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Anti-MOG Z12 variable light chain (Fragment).
Name=Gm1502; Synonyms=anti-MOG kappa;
Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria, Euarchontoglires; Glires; Rodentia; Sciurognathi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    EMBO J. 5:1577-1587(1986).
EMBL; AJ416331; CAC94866.1;
PIR; G27887; G27887.
PISSP; P01834; IMIM.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        response to a defined antigenic hemagglutinin.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Chernajovsky Y.;
Submitted (OCT-2001) to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sembi P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Muridae; Murinae; Mus.
NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Caton A.J., Brownlee G.G., Staudt L.M., "Structural and functional implications
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PubMed=242733
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRAIN=BALB/C;
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                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
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Created)
Last sequence update)
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Pred. No. 3
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Imunoglobulin gamma-3 kappa chain Mus musculus (Mouse).

precursor (Fragment).

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Q8X1F1 MOUSE
ID Q8K1F1 MOUSE
ID Q8K1F1 MOUSE
ID Q8K1F1 MOUSE
DT 01-0CT-2
DT 01-0CT-2
DT 01-MAR 2
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Best Local S
Matches 89
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SIGNAL
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NON TER
SEQUENCE
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EMBL; D14629; BAA03482.1; -; mRNA.
HSSP; P01679; 2FBJ.
SMR; Q811C3; 23-131.
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STRAIN-MRI/MpJ-lpr/lpr; TISSUS=Spleen;

MEDLINE=93156722; PubMed=8429833; DOI=10.1016/0161-5890(93)90089-T;

Takahashi S., Ith J., Nose M., Ono M., Yamamoto T., Kyogoku M.;

"Cloning and cDNA sequence analysis of nephritogenic monoclonal antibodies derived from an MRL/lpr lupus mouse.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Bukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;
Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   QBKIF1 MOUSE PRELIMINARY; PRT; 114 AA.
QBKIF1,
QBKIF1,
QBKIF1,
QBKIF1,
Q1-OCT-2002 (TrEMBLrel. 22, Created)
Q1-OCT-2002 (TrEMBLrel. 22, Last sequence update)
Q1-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Anti-VIPase light chain variable region (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ensembl; ENSMUSG00000058987; Mus musculus.
InterPro; IPR0079110; Ig-1ke.
InterPro; IPR003956; Ig_v.
SMART; SM00406; IGv; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUCLEOTIDE SEQUENCE.

NUCLEOTIDE SEQUENCE.

STRAIN-BALB/c; TISSUE=Hyperimmunized spleen;

Zhou Y.-X., Taguchi H., Planque S., Karle S.,

Zhou Y.-X., Taguchi H., Planque S., Karle S.,

Submitted (MAY-2002) to the EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mus musculus (Mouse).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PROSITE, PS50835; IG_LIKE; 1.
        NUCLEOTIDE SEQUENCE.

MEDILINE=92381444; PubMed=1512540; DOI=10.1084/jem.176.3.761;

Tillman D.M., Jou N.T., Hill R.J., Marion T.N.;

"Both IgM and IgG anti-DNA antibodies are the products of clonally selective B cell stimulation in (NZB x NZW)F1 mice.";

J. Exp. Med. 176:761-779(1992).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                               nonmutated germ-line genes.";
Proc. Natl. Acad. Sci. U.S.A. 86:4624-4628(1989)
                                                                                                                                                                                                                                                                                                                                                             Baccala R., Quang T.V., Gilbert M., Ternynck T., Avrameas
"Two murine natural polyreactive autoantibodies are encode
                                                                                                                                                                                                                                                                                                                                                                                                                                               PubMed=2499887;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUCLEOTIDE SEQUENCE.
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131 AA;
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Pred. No. 2.1e-36;
B; Mismatches 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                immunoglobulin gamma-3 kappa
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Best Local
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PIR; A33933; A33933.
PIR; PH1058; PH1058.
HSSP; PO1837; 25C8.
SMR; Q8K1F1; 2-114.
Ensembl; ENSMUSG00000059896; Mus musculus.
InterPro; IPR007110; Ig-like.
InterPro; IPR0073596; Ig_v.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 1.
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SEQUENCE
           completed: November
  : 159.344 Becs
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                                                    VRFSGSGSGTSYSLTISRMEAEDAATYYCQQWSSNPLTFGSGTKLELKR
                                          ARFSGSGSGTSYSLTISSMEAEDAATYYCQQYHSYPRTFGGGTKLEIKR
                                                                               DIVLTOSPAIMSASPGEKVTMTCRASSSVSSSYLHWYQQXSGASPKLWIYSTSNLASGVP
                                                                                           114 AA;
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12163 MW;
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            21,
            2005, 12:04:05
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Pred. No. 4
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Perfect score:
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 200000000
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         Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.
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1: /cgn2_6/ptodata/1/iaa/5_COMB.pep:*

2: /cgn2_6/ptodata/1/iaa/6_COMB.pep:*

3: /cgn2_6/ptodata/1/iaa/H_COMB.pep:*

4: /cgn2_6/ptodata/1/iaa/RE_COMB.pep:*

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         GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
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    US-08-553-497A-20
US-08-553-497A-28
US-07-646-221-26
US-07-634-278-58
US-08-477-728-58
US-08-487-200-58
US-08-487-200-58
US-08-487-200-51
US-08-477-728-31
US-08-477-728-31
US-08-477-728-31
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US-08-487-200-31
US-08-487-200-31
US-08-487-200-31
US-08-553-497A-26
US-08-553-497A-26
US-09-726-219A-243
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Sequence 20, Appl Sequence 28, Appl Sequence 28, Appl Sequence 58, Appl Sequence 58, Appl Sequence 58, Appl Sequence 58, Appl Sequence 31, Appl Sequence 31, Appl Sequence 31, Appl Sequence 31, Appl Sequence 24, Appl Sequence 24, Appl Sequence 27, Appl Sequence 27, Appl Sequence 27, Appl Sequence 28, Appl Sequence 21, Appl Sequence 21, Appl Sequence 21, Appl Sequence 251, Appl Sequence 251, Appl Sequence 251, Appl Sequence 251, Appl Sequence 253, Appl Sequence 254, Appl Sequence 254, Appl Sequence 254, Appl Sequence 255, 
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US-08-553-497A-20
; Sequence 20, Application US/08553497A
; Patent No. 5844093
; GENERAL INFORMATION:
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45	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30	29	28
474	474	475	475	475	475	475	475	475	475	475	475	475	475	475	475	477	477
85.7	85.7	85.9	85.9	85.9	85.9	85.9	85.9	85.9	85.9	85.9	85.9	85.9	85.9	85.9	85.9	86.3	86.3
108	108	553	553	553	553	270	255	255	246	246	128	128	108	108	107	238	235
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US-09-196-522-240	US-09-726-219A-240	US-09-102-716-16	US-09-364-088-16	US-09-188-082-16	US-08-661-052-16	US-08-652-507-2	US-09-618-869-8	US-09-553-498-8	US-08-469-658-57	US-08-469-486-57	PCT-US95-07302-2	US-08-619-491-2	US-09-196-522-241	US-09-726-219A-241	US-08-211-202-3	US-09-798-689-21	US-09-910-099-17
Sequence 240, App			•	Sequence 16, Appl	-		Sequence 8, Appli			•	-	Sequence 2, Appli	Sequence 241, App	Sequence 241, App	, A	7 2	17,

## ALIGNMENTS

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US-08-553-497A-20
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                                                                                        COMPUTER: ISM PC COMPARTIBLE
COMPUTER: ISM PC COMPARTIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/553,497A
FILING DATE: 17-NOV-1995
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO PCT/EP95/00978
FILING DATE: 16-MAR-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: EP 94104160.0
FILING DATE: 17-MAR-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: EP 94118970.6
FILING DATE: 02-DEC-1994
ATTORNEY/AGENT INFORMATION:
NAME: HAMLET-KING, DIANA
REGISTRATION NUMBER: MERCK 1726
REFERENCE/DOCKET NUMBER: MERCK 1726
TELEPHONE: 703-243-6333
TELEPHONE: 703-243-6333
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Best Local Similarity 92.6%;
Matches 100; Conservative
TELEPAX: 703-243-11.
INFORMATION FOR SEQ ID NO: 28:
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
1 FNGTH: 242 mino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
ADDRESSEE: MILLEN, WHITE, ZELANO & BRANIGAN, P.C STREET: 2200 CLARENDON BLVD. SUITE 1400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: ANTI-EGFR SINGLE-CHAIN FVS AND ANTI-EGFR TITLE OF INVENTION: ANTIBODIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: amino acid
                                                         TELEPHONE: 703-243-6410
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ZIP: 22201
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ROSELL, ELISABET
BLASCO, FRANCESC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             KETTLEBOROUGH, C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               JAUME
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MARY M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 JAUME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 506; DB 1;
Pred. No. 6.6e-42;
1; Mismatches 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 1; Length 244;
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US-07-946-421-26
                                                                                                                                                        US-07-946-421-26
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                                                                               Matches
                                                                                                 Best
                                                                                                                Query Match
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Best Local 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: Bendig, Mary M.
APPLICANT: Kettleborough,
APPLICANT: Saldanha, Jose
                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: WO PCT/EP
FILING DATE: 04-MAR-1992
PRIOR APPLICATION DATA:
APPLICATION UNMBER: EP 911933
APPLICATION NUMBER: EP 911933
ATTORNEY/AGENT INFORMATION:
NAME: Hamlet-King, Diana
REGISTRATION NUMBER: 33,302
REFERENCE/DOCKST NUMBER: Merc
TELECOMMUNICATION INFORMATION:
TELECTRANCE: 703-243-6333
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/946,421
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: 06-NOV-1
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Ploppy disk
                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 42
CORRESPONDENCE ADDRESS:
ADDRESSE: Millen, White, Zelano & Branigan,
ADDRESSEE: Millen, White, Zelano & Branigan,
STREET: 2200 Clarendon Boulevard, Suite 1400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: Humanized and Chimeric Monoclonal TITLE OF INVENTION: Antibodies
                                                                                                 Local
                                                                                                                                                                                          TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: Arlington
STATE: Virginia
COUNTRY: U.S.A.
                                                                                                                                                                            TOPOLOGY:
                                                                                                                                                                                                                                                                                         TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           137 DIELTOSPTIMSASPGEKVIMTCSDSSSVSYMYWYQQKTGSSPRLLIYDTSNLASGVPVR 196
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 197 FSGSGSGTSYSLTISRMEAEDAATYYCQQWSSYPLTFGAGTKLBIK 242
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         61 FSGSGSGTSYSLTISRMEAEDAATYYCQQWS&NPLTFGSGTKLELK 106
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   26
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Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    91.3%;
Similarity 92.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Application US/07946421
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                                                                                                                                                                            linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          06-NOV-1992
                                                                           90.8%; Score 502; DB 1; Length 128; 94.2%; Pred. No. 7.5e-42; ive 3; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO PCT/EP92/00480
                                                                                                                                                                                                                                                                                                                                                                                                                                             EP 911933892
                                                                                                                                                                                                                                                     26:
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86 SGSGTSYSLTISRMEAEDAATYYCQQWSSHIFTFGSGTKLEIK 128

SGSGTSYSLTISRMEAEDAATYYCQQWSSNPLTFGSGTKLELK 106

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US-07-634-278-58
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                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 07/590,274
FILING DATE: 28-SEP-1990
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/310,252
FILING DATE: 13-FEB-1989
PRIOR APPLICATION NUMBER: US 07/290,975
FILING DATE: 28-DEC-1988
ATTONNEY/AGENT INFORMATION:
NAME: Smith, William M
REGISTRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 11823-00260
TELEPHONE: (415) 386-2400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                       TELEPHONE: (415) 326-2400
TELEPAX: (415) 326-2422
INFORMATION FOR SEQ ID NO: 58:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/634,278
FILING DATE: 19-DEC-1990
CLASSIFICATION: 424
PRIOD ADDOX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PC COMPATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
                                                                                                                                                                                                                                  TOPOLOGY: linear MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: Palo Alto
STATE: California
                                                                                                                                                                                                                                                                                     TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY:
                                                                                                                                       Match 90.6%;
Local Similarity 93.2%;
es 96; Conservative 4
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                                 64
 64
                                                                                          4 LTQSPAIMSASPGEKVTTTCSASSSVSYMYWYQQKPGSSPRLLIYDTSNLASGVPVRFSG 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      94301
                      8GSGTSYSLTISRMEAEDAATYYCQQWSSNPLTFGSGTKLELK 106
GGSGTSYSLTISRMEAEDAATYYCQQWSTYPLTFGAGTKLELK 106
                                                                       LTOSPAINSASPGEKVIMTCSGSSSVSFMYWYQQRPGSSPRLLIYDTSNLASGVPVRFSG 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Application US/07634278
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SELICK, Harold E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CO, Man Sung
SCHNEIDER, William P.
LANDOLFI, Nicholas F.
COBLINGH, Kathleen L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             QUEEN, Cary L.
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                                                                                                                                                                                                                                                                  single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    113
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                                                                                                                                                                                                                                                                                                                                                                                                                 11823-002600
                                                                                                                                           4
                                                                                                                                           Score 501; DB 1; Length 106; Pred. No. 7.6e-42; 4; Mismatches 3; Indels
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                                                                                                                                             Gaps
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RESULT 5

RESULT 6 US-08-474-040-58

; Sequence 58, Application US/08474040 ; Patent No. 5693761

GENERAL INFORMATION:
APPLICANT: QUEEN, Cary L.
APPLICANT: CO, Man Sung

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US-08-477-728-58
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GENERAL INFORMATION:
                                                                                                                                         Matches
                                                                                                                                                                       Query Match
                                                                                                                                                          Best
                                                                                                                                                                                                                                                                                                                 TELEFAX: (415) 326-2422
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                     PILING DATE: 13-FEB-1989
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/290
PILING DATE: 28-DEC-1988
ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M
REGISTRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 1182
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 326-2400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/477,728
FILING DATE: 07-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PC COMPATING SYSTEM: PC-DOS/MS-DOS
SOFTMARE: Patentin Release #1.0.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US 07/590,274
PILING DATE: 28-SEP-1990
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/310,252
PILING DATE: 13-FEB-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: QUBEN, CATY L.
APPLICANT: SCHNEIDER, Will
APPLICANT: SELICK, Harolo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0
FILING DATE: 19-DEC-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: IMPROVED HUMANIZED IMMUNOGLOBLINS
                                                                                                                                                                                                                                 MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION:
                                                                                                                                                                                                                                                STRANDEDNESS:
TOPOLOGY: lin
                                                                                                                                                          Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE:
                                                                                                                                                                                                                                                                                                    LENGTH:
64
                                                                                                                                        96;
                                                                                             4 LTQSPAIMSASPGEKVTTTCSASSSVSYMYMYQQKPGSSPRLLIYDTSNLASGVPVRFSG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    94111
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                                                                                                                                                          Similarity
                                                                                                                                                                                                                                                                                     amino acid
                           SGSGTSYSLTISRMEAEDAATYYCQQWSSNPLTFGSGTKLELK 106
 SGSGTSYSLTISRMEAEDAATYYCQQWSTYPLTFGAGTKLELK 106
                                                                    LTQSPAIMSASPGEKVTMTCSGSSSSVSFMYWYQQRPGSSPRLLIYDTSNLASGVPVRFSG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Application US/08477728
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      E: Townsend and Townsend and Crew LLP Two Embarcadero Center, 8th Floor
                                                                                                                                                                                                                                                                                                    106 amino acids
                                                                                                                                        Conservative
                                                                                                                                                                                                                                                  linear
                                                                                                                                                                                                                           peptide
                                                                                                                                                                                                                                                               single
                                                                                                                                                          90.6%;
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                                                                                                                                         4; Mismatches
                                                                                                                                                          Score 501; DB 1;
Pred. No. 7.6e-42;
                                                                                                                                                                                                                                                                                                                                                                                                                    11823-002600
                                                                                                                                                                             Length 106;
                                                                                                                                             Indels
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                                                                                                                                                                                     US-08-487-200-58
                                                           Sequence 58, Application US/08487200
Patent No. 5693762
GENERAL INFORMATION:
APPLICANT: QUEEN, Cary L.
APPLICANT: CO, Man Sung
APPLICANT: SCHNEIDER, William P.
                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US 07/310,252
PRICE APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US 07/290,975
PILING DATE: 28-DEC-1988
ATTORNEY/AGENT INFORMATION:
NAME: 6Mith, William M
REGISTRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 11823-002600
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 326-2400
TELEPAX: (415) 326-2422
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: (415) 326-2422
INFORMATION FOR SEQ ID NO: 58:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: SCHNEIDER, William P.
APPLICANT: LANDOLFI, Nicholas F.
APPLICANT: COELINGH, Kathleen L.
APPLICANT: SELICK, Harold B.
TITLE OF INVENTION: IMPROVED HUMANIZED IMMUNOGLOBLINS
NUMBER OF SEQUENCES: 113
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PM PC COMPATION
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION DATA:
APPLICATION UMBER: US/08/474,040
FILING DATE: 07-UN-1995
CLASSIFICATION: 536
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
LENGTH: 106 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/634,278
FILING DATE: 19-DEC-1990
APPLICATION NUMBER: US.07/590,274
                       APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 28-SEP-1990
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: Palo Alto
STATE: California
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                                                                                                                                                                                                                                                        64 SGSGTSYSLTISRMEABDAATYYCQQWSSNPLTFGSGTKLELK 106
                                                                                                                                                                                                                                                                                                                                                                                                                                 96,
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                                                                                                                                                                                                                                                                                                                                                   LTQSPAIMSASPGEKVTMTCSGSSSVSFMYWYQQRPGSSPRLLIYDTSNLASGVPVRFSG
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CO, Man Sung
SCHMEIDER, William E
LANDOLFI, Nicholas E
COELLNGH, Kathleen I
SELICK, Harold E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                106 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  90.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                               4; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 501; DB 1; Length 106; Pred. No. 7.6e-42;
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US-08-484-537-58
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                                                                                                                                                                                              Sequence 58, Application US/08484537
Patent No. 6180370
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTMARE: DETERMINE Release #1.0, Version #1.
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/487,200
FILING DATE: 7-JUN-1995
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION UMBER: US 07/634,278
FILING DATE: 19-DEC-1990
PRIOR APPLICATION DATA:
APPLICATION UMBER: US 07/590,274
FILING DATE: 28-SEP-1990
PRIOR APPLICATION UMBER: US 07/590,274
FILING DATE: 28-SEP-1990
PRIOR APPLICATION UMBER: US 07/310,252
FILING DATE: 13-FEB-1989
PRIOR APPLICATION UMUMBER: US 07/290,975
FILING DATE: 13-FEB-1989
PRIOR APPLICATION UMUMBER: US 07/290,975
FILING DATE: 28-DEC-1988
ATTORNEY/AGENT INFORMATION:
NAME: Smith William M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: (415) 326-2422
INFORMATION FOR SEQ ID NO: :
SEQUENCE CHARACTERISTICS:
LENGTH: 106 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Smith, nate: 30,223
REGISTRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 11823-002610
REFERENCE/DOCKET NUMBER: 11823-002610
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION (415) 326-2400
TELECOMMUNICATION (415) 326-2422
                APPLICANT: QUEEN, Cary L.
APPLICANT: CO, Man Sung
APPLICANT: SCHNEIDER, William P.
APPLICANT: LANDOLFI, Nicholas F.
APPLICANT: COELINGH, Kathleen L.
APPLICANT: SELICK, Harold B.
TITLE OF INVENTION: INFROVED HUMANIZED IMMUNOGLOBLINS
NUMBER OF SEQUENCES: 113
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: IMPROVED HUMANIZED IMMUNOGLOBLINS NUMBER OF SEQUENCES: 113
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and trew
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: peptide
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET: 379 Lytton Avenue CITY: Palo Alto
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                h 90.6%;
Similarity 93.2%;
                                                                                                                                                                                                                                                                                                                                                           GGSGTSYSLTISRMEAEDAATYYCQQWSTYPLTFGAGTKLELK 106
                                                                                                                                                                                                                                                                                                                                                                                                        SGSGTSYSLTISRMEAEDAATYYCOOWSSNPLTFGSGTKLELK 106
                                                                                                                                                                                                                                                                                                                                                                                                                                                      LTOSPAIMSASPGEKVIMICSGSSSVSFMYWYQQRPGSSPRLLIYDISNLASGVPVRFSG 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        California
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Pred. No. 7.6e-42;
4; Mismatches 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 1; Length 106
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ADDRESSEE: Townsend STREET: 379 Lytton CITY: Palo Alto

Avenue

Townsend and Townsend Khourie and Crew

California

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity
Matches 96; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                      Sequence 31, Appl. Patent No. 553010
                                                                          GENERAL INFORMATION:

APPLICANT: QUEEN, CARY L.

APPLICANT: CO, Man Sung
APPLICANT: SCHNEIDER, William P.

APPLICANT: COELINGH, Kathleen L.

APPLICANT: SELICK, Harold E.

TITLE OF INVENTION: IMPROVED HUMANIZED IMMUNOGLOBLINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/07/634,278
APPLICATION NUMBER: US/07/590,274
APPLICATION NUMBER: US 07/590,274
FILING DATE: 28-SEP-1990
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/310,252
FILING DATE: 13-FEB-1989
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US 07/290,975
FILING DATE: 28-DEC-1988
ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M
REGISTRATION NUMBER: 30,223
REPERENCE/DOCKET NUMBER: 11823-0026
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 326-2400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (415) 326-2422
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn Release #1.0, Version #1.25
                                       NUMBER OF SEQUENCES: 1. CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STRANDEDNESS: sir
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/08/484,537
                        ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                    64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      4 LTQSPAIMSASPGEKVTMTCSGSSSVSFMYWYQQRPGSSPRLLIYDTSNLASGVPVRFSG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             amino acid
                                                                                                                                                                                                                                                                                                                                                                                                          SGSGTSYSLTISRMEAEDAATYYCQQWSSNPLTFGSGTKLELK 106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LTQSPAIMSASPGEKVTTTCSASSSVSYMYWYQQKPGSSPRLLIYDTSNLASGVPVRFSG
                                                                                                                                                                                                                                                                                                                                                                                    SGSGTSYSLTISRMEAEDAATYYCQQWSTYPLTFGAGTKLELK 106
                                                                                                                                                                                                                                                                          Application US/07634278
379 Lytton Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   90.6%;
ilarity 93.2%;
Conservative
                        Townsend and Townsend Khourie and Crew
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 501; DB 2; Length 106; Pred. No. 7.6e-42;
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US-08-477-728-31
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                                                                                                                                                                                                                                                                                                                                                                                                                                vequence 31, Application US/08477728
Patent No. 5585089
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PILING DATE: 13-FEB-1989
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/290,975
FILING DATE: 28-DEC-1988
ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M
REGISTRATION NUMBER: 30,223
REFERENCE/DOCKST NUMBER: 11823-002600
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 326-2400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEPAX: (415) 326-2422
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: QUEEN, Cary
APPLICANT: SCHNEIDER, W
APPLICANT: SELICK, Haro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SUFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/634,278 FILING DATE: 19-DEC-1990 CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatil
OPERATING SYSTEM: PC-DOS,
SOFTWARE: Patentin Relean
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION NUMBER: US/08/477,728
APPLICATION NUMBER: US/08/477,728
                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, 8th floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0
FILING DATE: 28-SEP-1990
                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                 PPLICANT: SELICK, Harold
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 128 amino acids TYPE: amino acid TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: US
                                                                                                                                                                                                CITY: Palo Alto
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Match 90.6%; Local Similarity 93.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            64 SGSGTSYSLTISRMEAEDAATYYCQQWSSNPLTFGSGTKLELK 106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            26 LTQSPAIMSASPGEKVTMTCSGSSSVSFMYMYQQRPGSSPRLLIYDTSNLASGVPVRFSG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                4 LTQSPAIMSASPGEKVTTTCSASSSVSYMYMYQQKPGSSPRLLIYDTSNLASGVPVRFSG 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    96; Conservative
                                                                                                                                                            9411:
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SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                 IMPROVED HUMANIZED IMMUNOGLOBLINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US 07/310,252
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                                                                                                                                                                                                                                                                                                                                                                              William P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 501; DB 1; Length 128; Pred. No. 9.5e-42; 4; Mismatches 3; Indels
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Gaps

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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07
FILING DATE: 19-DEC-1990

US 07/634,278

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RESULT 11
US-08-474-040-31
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 28-DEC-1988
ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M
REGISTRATION NUMBER: 30,223
REPERBNOE/DOCKET NUMBER: 11823-002600
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 326-2400
TELEFAX: (415) 326-2422
INFORMATION FOR SEQ ID NO: 31:
SEQUENCE CHARACTERISTICS:
LENGTH: 128 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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Best Local &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTMARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/474,040
FILING DATE: 07-UN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 19-DEC-1990
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/590,274
FILING DATE: 28-SEP-1990
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/310,252
      CLASSIFICATION: 536
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: US 07/634,278
                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: QUEEN, Cary L.
APPLICANT: CO, Man Sung
APPLICANT: SCHNEIDER, William P.
APPLICANT: LANDOLFI, Nicholas F.
APPLICANT: COELINGH, Kathleen L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PILING DATE: 13-FEB-1989
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0
                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES: 1
                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: SELICK, Harold E.
TITLE OF INVENTION: IMPROVED HUMANIZED IMMUNOGLOBLINS
                                                                                                                                                                                                                                                                                 STREET: 379 Lytton Avenue CITY: Palo Alto STATE: California
                                                                                                                                                                                                                                                                                                                                              ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             86 SGSGTSYSLTISRMEAEDAATYYCQQWSTYPLTFGAGTKLELK 128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       64 SGSGTSYSLTISRMEAEDAATYYCQQWSSNPLTFGSGTKLELK 106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         96;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4 LTQSPAIMSASPGEKVTTTCSASSSVSYMYWYQQKPGSSPRLLIYDTSNLASGVPVRFSG 63
                                                                                                                                                                                                                                               94301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  H: 128 amino acids
amino acid
OGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        l, Application US/08474040
5693761
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LTQSPAIMSASPGEKVTMTCSGSSSVSFMYWYQQRPGSSPRLLIYDTSNLASGVPVRFSG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                              Townsend and Townsend Khourie and Crev
19-DEC-1990
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 501; DB 1;
Pred. No. 9.5e-42;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:

APPLICANT: COUESN, Cary L.

APPLICANT: CO, Man Sung

APPLICANT: SCHNEIDER, William P.

APPLICANT: LANDOLFI, Nicholas F.

APPLICANT: COELINGH, Kathleen L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
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Best Local Similarity
                                                                                 FILING DATA: 424
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: US 07/6
APPLICATION NUMBER: US 07/6
APPLICATION DATE: 19-DEC-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: (415) 326-2422 INFORMATION FOR SEQ ID NO: 31:
                                                                                                                                                                         COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATE: PatentIn Release #1.0, Version #1.25
SOFTWARE: PatentIn NOTA:
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/487,200
FILING DATE: 7-UN-1995
                                                  PRIOR APPLICATION DATA:
APPLICATION UMBER:
                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: SELICK, HAROLD E.
TITLE OF INVENTION: IMPROVED HUMANIZED AMMUNOGLOBLINS
NUMBER OF SEQUENCES: 113
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
LENGTH: 128 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REGISTRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 1TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 326-2400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US 07/290,975
PILING DATE: 28-DEC-1988
ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M
FILING DATE: 28-SEP-PRIOR APPLICATION DATA: APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                              COUNTRY: US
ZIP: 94301
                                                                                                                                                                                                                                                                                                                                                                                                         STREET: 379 Lytton Avenue CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: FILING DATE: 28-SEF
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              86 SGSGTSYSLTISRMEAEDAATYYCQQWSTYPLTFGAGTKLELK 128
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                                                                                                                                                                                                                                                                                                                                                                                    California
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                                       UMBER: US 07/590,274
28-SEP-1990
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                                                                                                                         US 07/634,278
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    US 07/310,252
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FILING DATE: 13-FEB-1
PRIOR APPLICATION DATA:
APPLICATION NUMBER: UPPLICATION NUMBER:

13-FEB-1989

28-DEC-1988

US 07/290,975

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US-08-484-537-31
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/07/634,278
APPLICATION NUMBER: US 07/590,274
APPLICATION NUMBER: US 07/590,274
APPLICATION NUMBER: US 07/590,274
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/310,252
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/290,975
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M
REGISTRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 1182
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 326-2400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: QUEEN,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                            COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/484,537
FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: US
ZIP: 94301
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend Khourie and Crew
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: SELICK, HAROLD E.
TITLE OF INVENTION: IMPROVED HUMANIZED IMMUNOGLOBLINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        UMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 128 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: Palo Alto
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180370
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GGGGTSYSLTISRMEAEDAATYYCQQWSTYPLTFGAGTKLELK 128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 California
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        379 Lytton Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SCHNEIDER, William P. LANDOLFI, Nicholas F. COELINGH, Kathleen L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CO, Man Sung
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   QUEEN, Cary L.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 501; DB 1; Length 128; Pred. No. 9.5e-42;
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GENERAL IN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: (415) 326-24:
INFORMATION FOR SEQ ID NO:
                                                                                               APPLICATION NUMBER: WO PCT FILING DATE: 16-MAR-1995 PRIOR APPLICATION DATA: BP 941 APPLICATION NUMBER: BP 941 FILING DATE: 17-MAR-1994 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REGISTRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 11
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 326-2400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT:
TITLE OF IN
TITLE OF IN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                     SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                           APPLICATION NUMBER: EP 94118970.6 FILING DATE: 02-DEC-1994 ATTORNEY AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS
                                                                                                                                                                                                      PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES:
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Local Similarity 93.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEPHONE:
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                                                                                                                                                                                                                                     APPLICATION NUMBER: US/08/553,497A FILING DATE: 17-NOV-1995
                                                                                                                                                                                                                                                                                                                                                                                                                STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET:
                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: ARLINGTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH:
                                                                                                                                                                                                                        CLASSIFICATION:
                NAME: HAMLET-KING, DIANA REGISTRATION NUMBER: 33,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         26 LTQSPAIMSASPGEKVTMTCSGSSSVSFMYWYQQRPGSSPRLLIYDTSNLASGVPVRFSG 85
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 4 LTQSPAINSASPGEKVTTTCSASSSVSYMYWYQQKPGSSPRLLIYDTSNLASGVPVRFSG 63
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Application US/08553497A
                                                                                                                                                                                                                                                                                                                                                                                                                                                   3: MILLEN, WHITE, ZELANO & BRANIGAN, P.C. 2200 CLARENDON BLVD. SUITE 1400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MITJANS, FRANSESC
ROSELL, ELISABET
BLASCO, FRANCESC
PIULATS, JAUME
VENTION: ANTI-EGFR SINGLE-CHAIN FVS AND ANTI-EGFR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           KETTLEBOROUGH, C. A. BENDIG, MARY M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GUSSOW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            JAUME
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                                                                                                                                                                                    WO PCT/EP95/00978
                                                                                                                                   BP 94104160.0
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MERCK 1726
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 501; DB 2; Length 128; Pred. No. 9.5e-42;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3; Indels
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, MOLECULE TYPE: protein US-08-553-497A-26

TELEPHONE: 703-243-6333
TELEFAX: 703-243-6410
INFORMATION FOR SEQ ID NO: 26
SEQUENCE CHARACTERISTICS:
LENGTH: 242 amino acids
TYPE: amino acid
TOPOLOGY: linear

26:

TELECOMMUNICATION INFORMATION:

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COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/553,497A
FILING DATE: 17-NOV-1995
CLASSIFICATION DATA:
APPLICATION NUMBER: WO PCT/EP95/00978
FILING DATE: 16-MAR-1995
PRIOR APPLICATION NUMBER: WO PCT/EP95/00978
FILING DATE: 16-MAR-1995
PRIOR APPLICATION NUMBER: BP 94104160.0
FILING DATE: 17-MAR-1994
APPLICATION NUMBER: EP 94104160.0
FILING DATE: 17-MAR-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: EP 94104160.6
FILING DATE: 02-DEC-1994
ATTORNEY/AGENT INFORMATION:
NAME: HAMLET-KING, DIANA
REGISTRATION NUMBER: MERCK 1726
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-243-6333
TETERENY TO A 243-6333
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Best Local Similarity 91.5%;
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APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: MITJANS, FRANSESC
APPLICANT: BLASCO, FRANCESC
APPLICANT: BLASCO, FRANCESC
APPLICANT: PIULATS, JAUME
TITLE OF INVENTION: ANTI-EGFR SINGLE-CHAIN FVS AND ANTI-EGFR
TITLE OF INVENTION: ANTIBODIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSEE: MILLEN, WHITE, ZELANO & BRANIGAN, P.C.
STREET: 2200 CLARENDON BLVD. SUITE 1400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: ARLINGTON STATE: VA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: U
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          197 FSGSGSGTSYSLTISRMEAEDAATYYCQQWSSYPLTFGAGTKLEIK 242
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    137 DIELTQSPTIMSASPGEKVTMTCSDSSSVSYTYWYQQKTGSSPRLLIYDTSNLASGVPVR 196
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       61 FSGSGSGTSYSLTISRMEAEDAATYYCOOMSSNPLTFGSGTKLELK 106
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                                                                                                                                        Query Match
Best Local Similarity
Matches 98; Conserv
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LENGTH: 246 amino acids
                                                                                                                                                                                                                                  MOLECULE TYPE: protein
                                                                                                                                                                                                                                                 TYPE: amino TOPOLOGY: li
199
                                                                 139 DIBLTQSPTIMSASPGEKVTMTCSDSSSVSYMYWYQQKPGSSPRLLIYDTSNLASGVPVR
                                                                                                                                          98, Conservative
                             61 FSGSGSGTSYSLTISRMEAEDAATYYCQQWSSNP--LTFGSGTKLELK 106
                                                                                                      1 DTELTQSPAIMSASPGEKVTTTCSASSSVSYMYWYQQKPGSSPRLLIYDTSNLASGVPVR
FSGSGSGTSYSLTISRMEAEDAATYYCQQWSSYPPMYTFGGGTKLEIK 246
                                                                                                                                                                                                                                                   linear
                                                                                                                                                           89.7%; Score 496; DB 1; Length 246; 90.7%; Pred. No. 6.3e-41;
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                                                                                                                                          1; Mismatches
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Search completed: November 21, 2005, 12:07:36
Job time: 41.37 secs

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Maximum DB seq length: 200000000
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Perfect score:
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Maximum Match 100%
Listing first 45 summaries
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 pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               and is derived by analysis of the total
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Gapop 10.0 , Gapext 0.5
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94.8
94.0
93.7
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Match Length
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 Published Applications AA Main:*

1: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*

2: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

3: /cgn2_6/ptodata/1/pubpaa/US09_PUBCOMB.pep:*

4: /cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*

5: /cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.pep:*

5: /cgn2_6/ptodata/1/pubpaa/US11_PUBCOMB.pep:*
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Copyright (c) 1993 - 2005 Compugen Ltd.
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US-10-632-706-87
US-09-144-886-78
US-09-144-886-90
US-10-632-706-88
US-10-344-779-2
US-10-738-809-2
US-10-895-135-46
US-09-144-886-75
US-10-632-706-72
US-10-632-706-72
US-10-632-706-72
US-10-468-370-643
US-10-468-370-643
US-10-4895-135-47
US-10-895-135-47
US-10-10-895-135-47
US-10-112-788-9
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                                       Sequence 87, Appl
Sequence 90, Appl
Sequence 90, Appl
Sequence 75, Appl
Sequence 2, Appl
Sequence 2, Appl
Sequence 2, Appl
Sequence 46, Appl
Sequence 75, Appl
Sequence 72, Appl
Sequence 17, Appl
Sequence 18, Appl
Sequence 21, Appl
Sequence 21, Appl
Sequence 24, Appl
Sequence 24, Appl
Sequence 24, Appl
Sequence 24, Appl
Sequence 2, Appl
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 Sequence
Sequence
Sequence
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Sequence 90, Application US/09144886
Patent NO. US20020155114A1

GENERAL INPORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
TITILE OF INVENTION: Botulinum Neurotoxins
FILE REFERENCE: 2500.117USO
CURRENT EPLICATION NUMBER: US/09/144,886
CURRENT FILING DATE: 1998-08-31
NUMBER OF SEG ID NOS: 98
SOFTWARE: Patentin Ver. 2.0
SEG ID NO 90
RESULT 4

US-10-632-706-88

| Sequence 88, Application US/10632706
| Publication No. US20040175385A1
| GENERAL INFORMATION:
| APPLICANT: MARKS, JAMES D.
| APPLICANT: AMERSDORFER, PETER
| APPLICANT: AMERSDORFER, PETER
| TITLE OF INVENTION: THERAPBUTIC MONOCLONAL ANTIBODIES THAT NEUTRALIZE BOTULINUM
| TITLE OF INVENTION: NEUTOTOXINS
| FILE REFERENCE: 407T-895120US
| CURRENT APPLICATION NUMBER: US/10/632,706
| CURRENT FILING DATE: 2003-08-01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; TYPE: PRT; ORGANISM: Artificial Sequence; ORGANISM: Artificial Sequence; PEATURE; OTHER INFORMATION: Description of Artificial Sequence: BoNT/A clone; OTHER INFORMATION: C1 region VL epitope 1
US-09-144-886-78
                                                                                                                                                                                                                                                                                                                                                             5
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US-09-144-886-90
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Best Local &
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NUMBER OF SEQ ID NOS: 98
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 78
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 94.0%;
Best Local Similarity 94.4%;
Matches 101; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION: Description of Artificial Sequence: BONT/A clone OTHER INFORMATION: 3F4 region VL epitope 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 107
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 520; DB 3; Length 107; Pred. No. 6.2e-39; Indels 1; Mismatches 5; Indels
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Pred. No. 2.7e-39;
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RESULT 6 US-10-344-779-2

Sequence 2, Application US/10344779
Publication No. US20030211106A1
GENERAL INFORMATION:
APPLICANT: SWEET, RAYMOND W.
APPLICANT: TORNSTTA, MARK A.

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PRIOR APPLICATION NUMBER: US 60/400,721
PRIOR PILING DATE: 2002-08-01
PRIOR PPLICATION NUMBER: US 09/144,806
PRIOR FILING DATE: 1998-08-31
NUMBER OF SEQ ID NOS: 278
SOFTWARE: Patentin version 3.2
SEQ ID NO 88
LENGTH: 109
TYPE: PRI
ORGANISM: Artificial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-10-632-706-75
Sequence 75, Application US/10632706
Publication No. US20040175385A1
GENERAL INFORMATION:
APPLICANT: MARKS, JAMES D.
APPLICANT: AMERSDORFER, PETER
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US-10-632-706-88
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                                                                                                                                                                                                                                                                                                                                                                    SEQ ID NO 75
LENGTH: 107
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Best Local Similarity 94.4%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: AMERSDORFER, PETTER
TITLE OF INVENTION: THERAPEUTE MONOCLONAL ANTIBODIES THAT NEUTRALIZE BOTULINUM
TITLE OF INVENTION: NEUTROTOXINS
TILE REFERENCE: 4077-895120US
CURRENT FILLE CATION NUMBER: US/10/632,706
CURRENT FILLING DATE: 2003-08-01
PRIOR APPLICATION NUMBER: US 60/400,721
PRIOR APPLICATION NUMBER: US 60/144,806
PRIOR FILLING DATE: 1908-08-31
                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS: 278
SOFTWARE: PatentIn version 3.2
                                                                                                                                                                                                                                                                                                                           TYPE: PRT
ORGANISM: Artificial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
                                                                                                                                                                                                                                                                       FEATURE:
OTHER INFORMATION: single chain antibody
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
61
                        61 FSGSGSGTSYSLTISRMEAEDAATYYCQQWS&NPLTFGSGTKLELKR 107
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                                                                                                               1 DTELTQSPAIMSASPGEKVTTTCSASSSVSYMYWYQQKPGSSPRLLIYDTSNLASGVPVR
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  FSGSGSGTSYSLTISRMEAEDAATYYCQQWSSYPLTFGAGTKLELKR 107
                                                                                    DIELTQSPAIMSASPGEKVIMTCSASSSVSYMYWYQQKPGSSPRLLIYDTSNLASQVPVR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        VRFSGSGSGTSYSLTISRMEABDAATYYCQQWSSNPLTFGSGTKLELKR
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                                                                                                                                                                          1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1; Mismatches
                                                                                                                                                                                                 Score 516; DB 4; Length 107; Pred. No. 1.4e-38;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 518; DB 4;
Pred. No. 9.5e-39;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 4; Length 109;
                                                                                                                                                                            Indels
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                                                                                          60
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; LENGTH: 129 ; TYPE: PRT ; ORGANISM: Mus musculus US-10-344-779-2

Query Match Best Local Similarity

93.3%;

Matches

100;

Conservative

2; Mismatches

Score 516; DB 4; Length 129; Pred. No. 1.7e-38; 2; Mismatches 2; Indels

0;

Gaps

NUMBER OF SEQ ID NOS: 18 SOFTWARE: PastSEQ for Windows Version 4.0 SEQ ID NO 2

FILE REFERENCE: GP50033

CURRENT APPLICATION NUMBER: US/10/344,779

CURRENT FILING DATE: 2003-02-17

PRIOR APPLICATION NUMBER: PCT/US01/26161

PRIOR FILING DATE: 2001-08-21

PRIOR APPLICATION NUMBER: US 60/226,524

PRIOR FILING DATE: 2000-08-21

PRIOR APPLICATION NUMBER: US 60/230,639

PRIOR APPLICATION NUMBER: US 60/230,639

PRIOR FILING DATE: 2000-09-07

PPLICANT: TRUNEH, ALEMSEGED

ITLE OF INVENTION: ANTI-RANK LIGAND MONOCLONAL ANTIBODIES

ITLE OF INVENTION: USEFUL IN TREATMENT OF BANK OF THE BEFORE THE B

RANK LIGAND MEDIATED DISORDERS

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4 LTQSPAIMSASPGEKVTMTCSASSSVSYMYWYQQKPGSSPRLLIYDTSNLASGVPVRFSG

63

SGSGTSYSLTISRMEAEDAATYYCQQWSSNPLTFGSGTKLELKR 107 SGSGTSYSLTISRMEAEDAATYYCQQWSNFPLTFGAGTKLELKR 107

LTQSPAIMSASPGEKVTTTCSASSSVSYMYWYQQKPGSSPRLLIYDTSNLASGVPVRFSG 63

64 64

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                                                                                                                                                                                               ; ORGANISM: Mus
US-10-738-809-2
                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/10/738,809
CURRENT FILING DATE: 2003-12-17
PRIOR APPLICATION NUMBER: US/10/344,779
PRIOR PILING DATE: 2003-02-19
PRIOR PILING DATE: 2003-02-19
PRIOR PILING DATE: 2001-08-21
PRIOR APPLICATION NUMBER: US 60/226,524
PRIOR PILING DATE: 2000-08-21
PRIOR PILING DATE: 2000-08-21
PRIOR PILING DATE: 2000-09-07
PRIOR PILING DATE: 2000-09-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: SWEET, I
APPLICANT: TORNETT
APPLICANT: TRUNEH
APPLICANT: WATTAM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 2, Application US/10738809
Publication No. US20040171117A1
                                                                                                                 Matches
                                                                                                                                   Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                           SEQ ID NO 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: TRUNEH, ALEMSEGED
APPLICANT: WATTAM, TREVOR A.
TITLE OF INVENTION: ANTI-RANK LIGAND MONOCLONAL ANTIBODIES
TITLE OF INVENTION: USEFUL IN TREATMENT OF RANK LIGAND MEDIATED DISORDERS
THERE REFERENCE: GP50033
                                                                                                                                                                                                                                                                                              NUMBER OF SEQ ID NOS: 18 SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                    LENGTH: 129
TYPE: PRT
64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TORNETTA, MARK A.
TRUNEH, ALEMSE
WATTAM
                                                           LTQSPAIMSASPGEKVTTTCSASSSVSYMYMYQQKPGSSPRLLIYDTSNLASGVPVRFSG 63
SGSGTSYSLTISRMEAEDAATYYCQQWSSNPLTFGSGTKLELKR 107
                                                                                                                   Conservative
                                                                                                                                                                                                                  musculus
                                                                                                                                 93.3%;
                                                                                                             Score 516; DB 4; Li
Pred. No. 1.7e-38;
Pred. matches 2;
                                                                                                                                                      Length 129;
                                                                                                                   Indels
                                                                                                                   0
                                                                                                                   Gaps
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US-10-895-135-46
; Sequence 46, Application US/10895135
; Publication No. US20050123549A1
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CURRENT APPLICATION NUMBER: US/10/895,135
CURRENT FILING DATE: 2004-07-21
PRIOR APPLICATION NUMBER: 60/488,447
PRIOR FILING DATE: 2003-07-21
NUMBER OF SEQ ID NOS: 63
SOFTWARE: PATENTIN VETSION 3.2
SEQ ID NO 46
SEQ ID NO 46
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 75, Application US/09144886 Patent No. US20020155114A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION
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                                                                                                                                                                                                                                                                                                                                   SEQ ID NO 75
                                                                                                                    Matches
                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Marks, James D
APPLICANT: Amersdorfer, Peter
APPLICANT: Amersdorfer, Peter
TITLE OF INVENTION: Therapeutic Monoclonal Antibodies That Neutralize
TITLE OF INVENTION: Botulinum Neurotoxins
FILE REFERENCE: 2500.117USO
CURRENT APPLICATION NUMBER: US/09/144,886
CURRENT FILING DATE: 1998-08-31
NUMBER OF SEQ ID NOS: 98
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: ImmunoGen, APPLICANT: PAYNE, Gi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: PAYNE, Gillian
APPLICANT: CHUN, Philip
APPLICANT: TAVARES, Daniel
TITLE OF INVENTION: A CA6 ANTIGEN-SPECIFIC
TITLE OF INVENTION: THE SAME
                                                                                                                                                                                                              FEATURE: OTHER INFORMATION: Description of Artificial Sequence: BONT/A clone OTHER INFORMATION: C15 region VL epitope 1
                                                                                                                                                                                                                                                                            ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                LENGTH: 107
TYPE: PRT
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61 FSGSGSGTSYSLTISRMEAEDAATYYCQQWSSNPLTFGSGTKLELKR 107
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                                                                   1 DTBLTQSPAIMSASPGEKVTTTCSASSSVSYMYWYQQKPGSSPRLLIYDTSNLASGVPVR 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         93.1%;
Similarity 91.6%;
                                                                                                                h 92.9%; Score 514; DB 3;
Similarity 92.5%; Pred. No. 2.1e-38;
99; Conservative 3; Mismatches 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DTELTOSPAINSASPGEKVTTTCSASSSVSYMYWYQQKPGSSPRLLIYDTSNLASGVPVR 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FSGSGSGTSYSLTISRMEAEDAATYYCQQWSTYPLTFGAGTKLELKR 107
                                        DIELTOSPAIMSASPGEKVIMTCSASSSVSHMYWYQQKPGSSPRLLIYDTSNLASGVPIR 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 515; DB 5;
Pred. No. 1.7e-38;
6; MismatcHes 3
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                                                                                                                      5
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                                                                                                                                                         Length 107,
                                                                                                                      Indels
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FILE REFERENCE: MER 118
CURRENT APPLICATION NUMBER: US/10/468,370
CURRENT FILING DATE: 2003-08-19
PRIOR APPLICATION NUMBER: EP 01103955.9
PRIOR PILING DATE: 2001-02-19
PRIOR PILING DATE: 2001-02-19
PRIOR FILING DATE: 2001-04-05
PRIOR FILING DATE: 2001-04-05
PRIOR APPLICATION NUMBER: PCT/EP02/01690
PRIOR APPLICATION NUMBER: PCT/EP02/01690
PRIOR PILING DATE: 2002-02-18
NUMBER OF SEQ ID NOS: 689
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 643
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; TYPE: PRT
; ORGANISM: Artificial
; FEATURE:
; OTHER INFORMATION: single chain antibody
US-10-632-706-72
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APPLICANT: MARKSDOFFER, PETER
APPLICANT: AMERSDOFFER, PONOCLONAL ANTIBODIES THAT NEUTRALIZE BOTULINUM
TITLE OF INVENTION: THERAPEUTIC MONOCLONAL ANTIBODIES THAT NEUTRALIZE BOTULINUM
TITLE OF INVENTION: NEUROTOXINS
FILE REFERENCE: 407T-895120US
CURRENT APPLICATION NUMBER: US/10/632,706
CURRENT APPLICATION NUMBER: US/60/400,721
PRIOR APPLICATION NUMBER: US/60/400,721
PRIOR APPLICATION NUMBER: US/60/400,721
PRIOR FILING DATE: 2002-08-01
PRIOR APPLICATION NUMBER: US/60/400,721
PRIOR FILING DATE: 1998-08-31
NUMBER OF SEG ID NO5: 278
SOFTWARE: PATENTIN VERBION 3.2
SEG ID NO 72
LENGTH: 107
TWORLS.
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                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: Gillies, Stephen
APPLICANT: Carr, Francis J.
APPLICANT: Jones, Tim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 643, Application US/10468370 Publication No. US20040082039A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
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Best Local Similarity
                                                                                                                                                                                                                                                                                                 APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                       APPLICANT: Way, Jeffrey
TITLE OF INVENTION: ARTIFICIAL PROTEINS WITH REDUCED
TITLE OF INVENTION: IMMUNOGENICITY
                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
LENGTH:
NO 643
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            61 FSGSGSGTSYSLTISRMEAEDSATYYCQQWSSYPFTFGSGTKLELKR 107
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                                                                                                                                                                                                                                                                                                                                                         Hamilton, Anita
Williams, Stephen
Hanlon, Marian
                                                                                                                                                                                                                                                                                                                                                                                                                    Jones, Tim
Carter, Graham
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                                                                                                                                                                                                                                                                                                                    Watkins, John
Baker, Matthew
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US-10-389-155-17
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US-10-468-528-2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
CORRESPONDENCE ADDRESS:
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Sequence 17, Application US/10389155 Publication No. US20030229208A1 GENERAL INFORMATION:
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LENGTH; 106
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 2, Application US/10468528 Publication No. US20040096442A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILE REFERENCE: MER-116
CURRENT APPLICATION NUMBER: US/10/468,528
CURRENT FILING DATE: 2003-08-19
PRIOR APPLICATION NUMBER: EP 01103954.2
PRIOR FILING DATE: 2001-02-19
PRIOR APPLICATION NUMBER: PCT/EP02/01687
PRIOR FILING DATE: 2002-02-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: MODIFIED ANTI-EGFR ANTIBODIES TITLE OF INVENTION: REDUCED IMMUNOGENICITY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Carter, Graham APPLICANT: Jones, Tim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Carr, Francis J. APPLICANT: Carter, Graham
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Wililams, Stephen
APPLICANT: Hamilton, Anita
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: PRT
ORGANISM: Mus Musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER INFORMATION: Antibody fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Match 90.8%;
Local Similarity 94.2%;
TITLE OF INVENTION: Improved Humanized Immunoglobulins NUMBER OF SEQUENCES: 100
                                                                                                                                                           APPLICANT: Queen, Cary L.
                                                                                                                                                                                                                                                                                                                                                                           64 SGSGTSYSLTISRMEAEDAATYYCQQWSSNPLTFGSGTKLELK 106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4 LTQSPAIMSASPGEKVTTTCSASSSVSYMYWYQQKPGSSPRLLIYDTSNLASGVPVRFSG 63
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                                      Co, Man Sung
Schneider, William P.
Landolfi, Nicholas F.
Coelingh, Kathleen L.
Selick, Harold E.
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Pred. No. 2.5e-37;
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; TOPOLOGY: linear; MOLECULE TYPE; protein; SEQUENCE DESCRIPTION: SEQ ID NO: 17: US-10-389-155-17
                                                                                                                                                                                                                                                RESULT 14
US-10-389-417-17
; Sequence 17, Application US/10389417
; Publication No. US20040049014A1
; GENERAL INFORMATION:
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SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION NUMBER: US/10/389,155
FILING DATE: 13-Mar-2003
PRIOR APPLICATION NUMBER: US/09/325,000
FILING DATE: 01-JUN-1999
APPLICATION NUMBER: US 07/290,975
FILING DATE: 28-DEC-1988
APPLICATION NUMBER: US 07/310,252
PILING DATE: 13-FEB-1989
APPLICATION NUMBER: US 07/590,274
PILING DATE: 28-SEP-1990
APPLICATION NUMBER: US 07/590,274
PILING DATE: 28-SEP-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Match 90.6%;
Local Similarity 93.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US 08/484,537 FILING DATE: 07-JUN-1995 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Smith, William M.
REGISTRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 01:
TELECOMMUNICATION INFORMATION:
                                                                Schneider, William P.
Landolfi, Nicholas F.
Coelingh, Kathleen L:
Selick, Harold E.
TITLE OF INVENTION: Improved Humanized Immunoglobulins
NUMBER OF SEQUENCES: 100
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                               APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                             64
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                                                                                                                                                                                                                                                                                                                                                                                                             SGSGTSYSLTISRMEAEDAATYYCQQWSSNPLTFGSGTKLELK 106
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 106 amino acids
TYPE: amino acid
STRANDEDNESS: <Unknown>
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TELEPAX: (415) 576-0300
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                                                                                                                                   Co, Man Sung
Schneider, William P.
Landolfi, Nicholas F.
Coelingh, Kathleen L.
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Pred. No. 3.1e-37;
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  GENERAL INFORMATION:

APPLICANT: Queen, Cary
APPLICANT: Co, Man Sung
APPLICANT: Schneider, William
APPLICANT: Landolfi, Nicholas
APPLICANT: Landolfi, Nicholas
APPLICANT: Landolfi, Nicholas
APPLICANT: Selick, Harold
TITLE OF INVENTION: Improved Humanized Immunoglobulins
FILE REFERENCE: 05882.0078.CNUS01
CURRENT APPLICATION NUMBER: US/10/452,357
CURRENT FILING DATE: 2003-05-30
PRIOR APPLICATION NUMBER: 09/718,993
PRIOR FILING DATE: 2000-11-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRANDEDNESS: «Unknown»

TOPOLOGY: linear

MOLECULE TYPE: protein

SEQUENCE DESCRIPTION: SEQ ID NO: 17:

US-10-389-417-17
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US-10-452-357-58
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Best Local S
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INFORMATION FOR SEQ ID NO: 17:
SEQUENCE CHARACTERISTICS:
LENGTH: 106 amino acids
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PILING DATE: 01-JUN-1999
APPLICATION NUMBER: US/09/325,000
PILING DATE: 01-JUN-1999
APPLICATION NUMBER: US 07/290,975
PILING DATE: 28-DEC-1988
APPLICATION NUMBER: US 07/310,252
PILING DATE: 13-FEB-1989
APPLICATION NUMBER: US 07/590,274
PILING DATE: 28-SEP-1990
APPLICATION NUMBER: US 07/634,278
PILING DATE: 19-DEC-1990
APPLICATION NUMBER: US 08/634,278
PILING DATE: 19-DEC-1990
APPLICATION NUMBER: US 08/484,537
PILING DATE: 07-JUN-1995
ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M.
REGISTRATION NUMBER: 30,223
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: 011823-002650US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION DATA:
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1 Similarity 93.2%;
96; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn Release #1.0, Version #1.30
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Pred. No. 3.1e-37;
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PRIOR APPLICATION NUMBER: 09/487,200

PRIOR PILING DATE: 1995-06-07

PRIOR PPLICATION UNMBER: 07/634,278

PRIOR PILING DATE: 1990-12-19

PRIOR PELLING DATE: 1990-12-19

PRIOR PILING DATE: 1990-02-13

PRIOR PILING DATE
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Scoring table:
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Maximum Match 100%
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 Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being prir and is derived by analysis of the total score distribution.
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1: /cgn2 6/ptodata/2/pubpaa/US10 NEW_PUB.pep:*

2: /cgn2_6/ptodata/2/pubpaa/US06_NEW_PUB.pep:*

3: /cgn2_6/ptodata/2/pubpaa/US07 NEW_PUB.pep:*

4: /cgn2_6/ptodata/2/pubpaa/US08_NEW_PUB.pep:*

5: /cgn2_6/ptodata/2/pubpaa/US09_NEW_PUB.pep:*

6: /cgn2_6/ptodata/2/pubpaa/US10_NEW_PUB.pep:*

7: /cgn2_6/ptodata/2/pubpaa/US11_NEW_PUB.pep:*

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Match
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US-11-174-186-16

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US-11-721-763-3-19

US-10-495-664-3

US-10-721-763-27

US-10-648-816-2

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                      Sequence 36, Appli
Sequence 1, Appli
Sequence 8, Appli
Sequence 42, Appli
Sequence 9, Appli
Sequence 9, Appli
Sequence 16, Appli
Sequence 16, Appli
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Sequence 19, Appli
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10-648-81 10-648-81 10-648-81 10-648-81 11-077-97 10-512-18 10-512-18 10-512-18 10-512-18 10-512-18 10-639-79 11-144-24 10-639-79 11-174-18 11-174-18 11-174-18 11-174-18 11-18-24 11-18-24 11-18-24 11-15-18	329	259	236	126	126	106	127	126	236	126	126	263	329	259	111	110	110	110	110	1
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	US-10-512-184-70	US-10-512-184-33	-11-144-248	-10-839-799	US-10-839-799-43	-11-174-186	-10-721-763	-10-839-799	-11-144-248	-10-839-799	10-839-799	-10-512-184	-10-512	-10-512-18	-11-077-97	-10-648-81	-10-	-10-	-10-648	
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4, Appli 6, Appli 8, Appli 11, Appli 11, Appli 11, Appli 29, Appli 29, Appli 48, Appli 48, Appli 48, Appli 48, Appli 48, Appli 48, Appli 48, Appli 48, Appli 53, Appli 54, Appli	App	App	Appl	Appl	Appl	Tddy	rddy	App	App	App	App	App	App	ddy	rade	Tdd	rada	rad	Tdd	

## ALIGNMENTS

RESULT 1 US-10-512-184-36

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RESULT 2
US-10-512-184-50
US-10-512-184-50
; Sequence 50, Application US/10512184
; Publication No. US20050244901A1
; GENERAL INFORMATION:
; APPLICANT: Fraunhofer Gesellschaft zu;
; APPLICANT: Fraunhofer Gesellschaft zu;
; TITLE OF INVENTION: Antibody fragment
; TITLE OF INVENTION: registance again
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; LENGTH: 248; TYPE: PRT
; ORGANISM: Artificial Sequence
; PEATURE;
OTHER INFORMATION: Description of Artificial Sequence: scFv SS2 with
; OTHER INFORMATION: specificity against Sclerotinia sclerotiorum;
; OTHER INFORMATION: originates from Mus musculus.
US-10-512-184-36
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Publication No. US20050244901A1
GENERAL INFORMATION:
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FILE REFERENCE: 3581.01US01
CURRENT APPLICATION NUMBER: US/10/512,184
CURRENT FILING DATE: 2004-10-22
NUMBER OF SEQ ID NOS: 72
SOPTWARE: Patentin Ver. 2.1
SEQ ID NO 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local (
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               APPLICANT: Fraunhofer Gesellschaft zur F"rderung der angewandten Forschung TITLE OF INVENTION: Antibodies, recombinant TITLE OF INVENTION: antibody fragments and fusions mediated plant disease TITLE OF INVENTION: resistance against fungi
CURRENT APPLICATION NUMBER: US/10/512,184
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1 Similarity 81.3%;
87; Conservative 7
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Sequence 1, Application US/11174186

Publication No. US20050244418A1

GENERAL INFORMATION:
APPLICANT: Gillies, Stephen
APPLICANT: Lo, Kin-Ming
APPLICANT: Qian, Xiugi
ITILE OF INVENTION: Recombinant Tumor Specific Antibody And Use These Pile REFERENCE: LEX-019

CURRENT APPLICATION NUMBER: US/11/174,186

CURRENT APPLICATION NUMBER: US 60/288,564

PRIOR APPLICATION NUMBER: US 60/288,564

PRIOR APPLICATION NUMBER: US 60/288,564

PRIOR FILING DATE: 2001-05-03

NUMBER OF SEQ ID NOS: 42

SOFTWARE: Patentin version 3.3

LENGTH. 106
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; NUMBER OF SEQ ID NOS: 72
; SOFTWARE: Patentin Ver. 2.1
; SEQ ID NO 50
; LENGTH: 615
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE: PRORMATION: Description of Artificial Sequence; fusion protein
; OTHER INFORMATION: comprising the leader peptide - chitinase - linker
; OTHER INFORMATION: - scFv SS2 - cmyc/H1s6.
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US-11-174-186-1
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Sequence 8, Application US/11174186
Publication No. US20050244418A1
GENERAL INFORMATION:
APPLICANT: Gillies, Stephen
APPLICANT: Lo, Kin-Ming
APPLICANT: Qian, Xiugi
TITLE OF INVENTION: Recombinant Tumor Specific Antibody And Use Thereof
FILE REFERENCE: LEX-019
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Best Local S
Matches 86
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Best Local Similarity 81.3
Matches 87; Conservative
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OTHER INFORMATION: KS VK mouse
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Publication No. US20050244413A1
GENERAL INFORMATION:
APPLICANT: Adolf, Guenther
APPLICANT: Heider, Karl-Heinz
APPLICANT: Heider, Karl-Heinz
TITLE OF INVENTION: Compositions and Methods for Treating Cancer using
TITLE OF INVENTION: Cytotoxic CD44 Antibody Immunoconjugates and
TITLE OF INVENTION: Chemotherapeutic Agents
FILE REFERENCE: 1/1383
CURRENT APPLICATION NUMBER: US/11/172,320
CURRENT FILING DATE: 2005-06-30
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LENGTH: 106
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CURRENT FILING DATE: 2005-07-01
PRIOR PPLICATION NUMBER: US 60/288,564
PRIOR FILING DATE: 2001-05-03
NUMBER OF SEQ ID NOS: 42
SOFTWARE: Patentin version 3.3
                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: US/10/645,215
PRIOR FILING DATE: 2003-08-21
PRIOR APPLICATION NUMBER: EP 02 018 686.2
PRIOR FILING DATE: August 21, 2002
PRIOR PILING DATE: August 21, 2002
PRIOR PILING DATE: August 26, 2002
NUMBER OF SEQ ID NOS: 9
                                                                                                                                                                                                                                                                         OTHER INFORMATION: Humanised Murine Antibody BIWA 4 Light Chain
                                                                                                                                                                                                                                                                                            ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                       TYPE: PRT
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ORGANISM: Artificial
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                                                                                                                                                                                                                                                                                                                                                         LENGTH: 213
                                                                                                                                                                                         Local
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SGSGTDFTLTISSLEPEDFAVYYCLQWSSNPLTFGGGTKVEIKR
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81.6%;
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Pred. No. 8e-22;
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RESULT 6
US-11-174-186-42
; Sequence 42, Application US/11174186

Publication No. US20050244418A1 GENERAL INFORMATION:

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; TYPE: PRT; ; CRATUISM: Artificial ; PEATURE; ; CTHER INFORMATION: light chain US-11-174-186-42
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RESULT 8
US-11-172-320-8
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TITLE OF INVENTION: Recombinant Tumor Specific Antibody And Use Thereof
FILE REFERENCE: LEX-019
CURRENT APPLICATION NUMBER: US/11/174,186
CURRENT FILING DATE: 2005-07-01
PRIOR APPLICATION NUMBER: US 60/288,564
PRIOR PILING DATE: 2001-05-03
NUMBER OF SEQ ID NOS: 42
SOFTWARE: Patentin version 3.3
SEQ ID NO 42
LENGTH: 213
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Best Local Similarity
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SOFTWARE: PatentIn version 3.3
SEQ ID NO 9
LENGTH: 106
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CURRENT APPLICATION NUMBER: US/11/174,186
CURRENT FILING DATE: 2005-07-01
PRIOR APPLICATION NUMBER: US 60/288,564
PRIOR FILING DATE: 2001-05-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Gillies, Stephen APPLICANT: Lo, Kin-Ming APPLICANT: Qian, Xiugi
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                                                                                                                                                                                                                                                                                                                                      FEATURE: OTHER INFORMATION: VK8 light chain
                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Artificial
                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: PRT
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Local Similarity 79.6%;
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Pred. No. 2.1e-21;
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Pred. No. 2.5e-21;
5; Mismatches 16
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                                                                                                                                                                 ; OTHER INFORMATION: Vk6 light chain US-11-174-186-7
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CURRENT FILING DATE: 2005-06-30
PRIOR APPLICATION NUMBER: US/11/645,215
PRIOR FILING DATE: 2003-08-21
PRIOR APPLICATION NUMBER: EP 02 018 686.2
PRIOR APPLICATION NUMBER: EP 02 018 686.2
PRIOR APPLICATION NUMBER: US 60/405,956
PRIOR APPLICATION NUMBER: US 60/405,956
PRIOR FILING DATE: August 26, 2002
NUMBER: US 60/405,956
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 7, Application US/11174186 Publication No. US20050244418A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQ ID NOS: 9
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 8
LENGTH: 213
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Adolf, Guenther
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                                                                                                     Query Match
Best Local :
                                                                                                                                                                                                                                                                                               SEQ ID NO 7
                                                                                                                                                                                                                                                                                                    APPLICANT: Gillies, Stephen
APPLICANT: Lo, Kin-Ming
APPLICANT: Olan, Xiugi
TITLE OF INVENTION: Recombinant Tumor Specific Antibody And Use Thereof
FILE REPERENCE: LEX-019
FILE REPERENCE: LEX-019
CURRENT APPLICATION NUMBER: US/11/174,186
CURRENT FILING DATE: 2005-07-01
PRIOR APPLICATION NUMBER: US 60/288,564
PRIOR FILING DATE: 2001-05-03
NUMBER OF SEQ ID NOS: 42
SOFTWARE: PatentIn version 3.3
                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Baum, Anke
APPLICANT: Heider, Karl-Heinz
TITLE OF INVENTION: Compositions and Methods
TITLE OF INVENTION: Cytotoxic CD44 Antibody
TITLE OF INVENTION: Chemotherapeutic Agents
FILE REFERENCE: 1/1383
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OTHER INFORMATION: Humanised Antibody BIWA 8 Light Chain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                               TYPE: PRT
ORGANISM: Artificial
                                                                                                                                                                                                           FEATURE:
                                                                                                                                                                                                                                                                       LENGTH: 106
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Similarity 73.1%;
                                                                                                        Similarity
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                        LTQSPAIMSASPGEKVTTTCSASSSVSYMYWYQQKPGSSPRLLIYDTSNLASGVPVRFSG 63
LTQSPATLSLSPGERVTLTCSASSSVSYMLWYQQKPGQAPKLLIFDTSNLASGIPARFSG
                                                                                    Conservative
                                                                                                     74.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; Score 416; DB 7; Length 213; ; Pred. No. d.5e-21; 13; Mismatches 15; Indels
                                                                                 Score 409; DB 7
Pred. No. 1e-20;
9; Mismatches
                                                                                                                            DB 7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       for Treating Cancer using
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                                                                                      17,
                                                                                                                       Length 106;
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Sequence 15, Application US/11174186

Publication No. US20050244418A1

GENERAL INFORMATION:
APPLICANT: Lo, Kin-Ming
APPLICANT: Lo, Kin-Ming
APPLICANT: Qian, Xiugi
TITLE OF INVENTION: Recombinant Tumor Specific Antibody And Use Thereof
FILE REFERENCE: LEX-019
FILE REFERENCE: LEX-019
CURRENT APPLICATION NUMBER: US/11/174,186
CURRENT APPLICATION NUMBER: US/67
FRIOR APPLICATION NUMBER: US/07-08
PRIOR APPLICATION NUMBER: US/07-08
SPINOR APPLICATION NUMBER: US 60/288,564
PRIOR APPLICATION NUMBER: US 60/288,564
PRIOR FILING DATE: 2001-05-03
NUMBER OF SEQ ID NOS: 42
SOFTWARE: PatentIn version 3.3
ISOCITAL 106
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US-11-174-186-16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Gillies, Stephen
APPLICANT: Lo, Kin Ming
APPLICANT: Qian, Xiugi
TITLB OF INVENTION: Recombinant Tumor Specific Antibody And Use Thereof
FILE REFERENCE: LEX-019
CURRENT APPLICATION NUMBER: US/11/174,186
CURRENT FILING DATE: 2005-07-01
PRIOR APPLICATION NUMBER: US 60/288,564
PRIOR APPLICATION NUMBER: US 60/288,564
PRIOR PRIOR DATE: 2001-05-03
NUMBER OF SEQ ID NOS: 42
SOFTWARE: Patentin version 3.3
SEQ ID NO 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 16, Application US/11174186 Publication No. US20050244418A1 GENERAL INFORMATION:
                                                                                    Matches
                                                                                                                        Query Match
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                                                                                                                                                                          FEATURE:
OTHER INFORMATION: KS de-immunized VK5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Artificial FEATURE:
                                                                                                                                                                                                                 ORGANISM: Artificial
                                                                                                                                                                                                                                           TYPE: PRT
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                                                                                                                                                                                                                                                             LENGTH:
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                                                                                                    Local Similarity
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                          4 LTQSPAIMSASPGEKVTTTCSASSSVSYMYMYQQKPGSSPRILLIYDTSNLASGVPVRFSG
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        LTQSPASLAVSPGQRATITCSASSSVSYMLWYQQKPGSSPKPWIYDTSNLASGFPARFSG 63
                                                                                  Conservative
                                                                                                  73.6%; Score 407; DB 7; Length 106; 74.8%; Pred. No. 1.4e-20;
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                                                                                10; Mismatches
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                                                                                  16; Indels
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US-11-174-186-10
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US-11-174-186-14
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                                                                                                                                                                                                                     SOFTWARE: PatentIn version 3.3
SQ ID NO 14
LENGTH: 106
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: Gillies, Stephen
                                                              Matches
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                                                                                                                                                                              ORGANISM: Artificial FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OTHER INFORMATION: KS VK veneered
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                                                                74;
  4 LTQSPAIMSASPGEKVTTTCSASSSVSYMYWYQQKPGSSPRLLIYDTSNLASGVPVRFSG 63
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                                                                Conservative
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Sequence 14, Application US/11174186
Publication No. US20050244418A1
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                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Gillies, Stephen
APPLICANT: Lo, Kin-Ming
APPLICANT: Qian, Xiugi
TITLE OF INVENTION: Recombinant Tumor Specific Antibody And Use Thereof
PILS REFERENCE: LEX-019
CURRENT APPLICATION NUMBER: US/11/174,186
CURRENT FILING DATE: 2005-07-01
PRIOR APPLICATION NUMBER: US 60/288,564
PRIOR FILING DATE: 2001-05-03
NUMBER OF SEQ ID NOS: 42
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APPLICANT: Qian, Xiugi
TITLE OF INVENTION: Recombinant Tumor Specific Antibody And Use Thereof
FILE REFERENCE: LEX-019
FILE REFERENCE: LEX-019
FILE REFERENCE: LEX-019
CURRENT APPLICATION NUMBER: US/11/174,186
CURRENT FILING DATE: 2005-07-01
PRIOR APPLICATION NUMBER: US 60/288,564
PRIOR FILING DATE: 2001-05-03
NUMBER OF SEQ ID NOS: 42
COMMENTS TRANSPORTED TO SECRET TO THE PRIOR PRIOR FILING DATE: 2001-05-03
NUMBER OF SEQ ID NOS: 42
COMMENTS TRANSPORTED TO THE PRIOR PRIOR PRIOR FILING DATE: 2001-05-03
NUMBER OF SEQ ID NOS: 42
                                                                                                                                                                          OTHER INFORMATION: KS de-immunized VK4
                               / Match 71.2%; Score 394; DB 7; Length 106, Local Similarity 71.8%; Pred. No. 8.7e-20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                64 SGSGTSYTLTINSLEAEDAATYYCHQRSGYPYTFGGGTKVEIK 106
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   71.2%; Score 394; DB 7; Length 106; 71.8%; Pred. No. 8.7e-20;
11; Mismatches 18; Indels
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4 LTQSPASLAVSPGQRATITCSASSSVSYMLWYQQKPGQPPKPWIFDTSNLASGFPARFSG 63

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RESULT 14
US-11-174-186-12
Sequence 12, Application US/11174186
; Publication No. US20050244418A1
; Publication No. US20050244418A1
; GENERAL INFORMATION:
; APPLICANT: Gillies, Stephen'
; APPLICANT: Lo, Kin-Ming
; APPLICANT: Olan, Xiugi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 15
US-11-174-186-13
Sequence 13, Application US/11174186
Publication No. US20050244418A1
GENERAL INFORMATION:
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US-11-174-186-12
                                                                                                                                            OTHER INFORMATION: KS-deimmunized VK3 US-11-174-186-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Qian, Xiugi
TITLE OF INVENTION: Recombinant Tumor Specific Antibody And Use Thereof
FILE REFERENCE: LEX-019
CURRENT APPLICATION NUMBER: US/11/174,186
CURRENT FILING DATE: 2005-07-01
PRIOR APPLICATION NUMBER: US 60/288,564
PRIOR FILING DATE: 2001-05-03
NUMBER OF SEQ ID NOS: 42
SOFTWARE: Patentin version 3.3
SEQ ID NO 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 70.9%; Score 392; DB 7; Length 106; Best Local Similarity 71.8%; Pred. No. 1.2e-19; Matches 74; Conservative 11; Mismatches 18; Indels
                                                                                                                                                                                                                                                                              PILE REFERENCE: LEX-019
CURRENT APPLICATION NUMBER: US/11/174,186
CURRENT PILING DATE: 2005-07-01
PRIOR APPLICATION NUMBER: US 60/288,564
PRIOR FILING DATE: 2001-05-03
NUMBER OF SEQ ID NOS: 42
SOFTWARE: Patentin version 3.3
SEQ ID NO 13
                                          Query Match 70.9%; Score 392; DB 7; Length 106; Best Local Similarity 71.8%; Pred. No. 1.2e-19; Matches 74; Conservative 11; Mismatches 18; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 106
TYPE: PRT
ORGANISM: Artificial
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Gillies, Stephen
APPLICANT: Lo, Kin-Ming
APPLICANT: Olan, Xingi
TITLE OF INVENTION: Recombinant Tumor Specific Antibody And Use Thereof
                                                                                                                                                                                      ORGANISM: Artificial FEATURE:
                                                                                                                                                                                                                                        LENGTH: 106
TYPE: PRT
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     4 LTQSPAIMSASPGEKVTTTCSASSSVSYMYWYQQKPGSSPRLLIYDTSNLASGVPVRFSG 63
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SGSGTSYSLTISRMEAEDAATYYCQQWSSNPLTFGSGTKLELK 106
                     63
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Search completed: November 21, 2005, 12:33:50 Job time : 2.05971 secs

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Minimum DB :
Maximum DB :
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Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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Copyright (c) 1993 - 2005 Compugen Ltd.
geneseqp2001s:*
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328.182 Million cell updates/sec
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

### SUMMARIES

24	23	22	21	20	19	18	17	16	15	14	13	12	11	10	9	8	7	0	ű	4	ω	N	1	No.	1
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AAE27810	ABG32682	ADO47747	AAB69655	AAR21287	ADO47720	AAB69673	AAR21304	AAR25958 .	AAR21306	AAR79871	AAR79870	ADR38691	ADT74993	AAR79867	ADR38670	ABB75616	AAR21305	ADU86571	ADU86565	ADW86126	ADR38673	ADR38685	ADR38686	ID	
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45	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30	29	28	27	26	25
469	469	469	469	469	469	469	470	470	470	471	471	471	471	471.5	471.5	472	473	473	475	475
83.9	83.9	83.9	83.9	83.9	83.9	83.9	84.1	84.1	84.1	84.3	84.3	84.3	84.3	84.3	84.3	84.4	84.6	84.6	85.0	85.0
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ADV14577	AAG66521	AAR21307	AAE22196	ADZ83490	ADZ64982	ADV66151	ADH59707	ADH59678	ADO16879	ADT75053	ADT75054	ADT75125	AAR21293	ADU86519	ADW86127	AAR79869	ADW80852	AAU72834	AAR79872	AAR27050
Adv14577	Aag66521	Aar21307	Aae22196	Adz83490	Adz64982	Adv66151	Adh59707	Adh59678	Ado16879	Adt75053	AGE 75054	Adt75125	Aar21293	Adu86519	Adw86127	Aar/9869	Adwansz	ABU/2834	ABT/98/2	Aar27050
VL (CD3) -V	Mouse ant	Murine VL	CD3-spec1	Non-deimm	Non-deimm	CD3-speci	Chimeric	CPAM4 VK		` ^	Light cha		m	orbourmmt	Novel cyc		Anci-Cus	AIRCI -NAGZ	ANCI-EGER	VL425 ant

### ALIGNMENTS

ADR38686 standard; peptide; 109 AA.

ADR38686;

02-DEC-2004 (first entry)

Mouse light chain variable region scFv seqid 88

antibacterial; antibody; botulinum neurotoxin type A; BONT/A; BONT/A neutralising epitope; anti-botulinum neurotoxin antibody; toxin neutralisation; botulinum neurotoxin poisoning; mouse; light chain variable region; single chain antibody.

Mus sp.

US2004175385-A1.

09-SEP-2004.

01-AUG-2003; 2003US-00632706

31-AUG-1998; 98US-00144886. 01-AUG-2002; 2002US-0400721P.

(REGC ) UNIV CALIFORNIA.

WPI; 2004-652009/63.

Marks JD,

Amersdorfer

P ;

New isolated antibody that neutralizes botulinum neurotoxin type A, useful for diagnosing botulism or for treating pathologies associated with botulinum neurotoxin poisoning.

Example 1; SEQ ID NO 88; 110pp; English.

RESULT 1
ADR3861
ID ADR37
XX ADR32
XX ADR32
XX ADR32
XX MOUS
DI MOUS
XX MUS
DI MOUS
XX MUS
DI MOUS
XX IIGH
XX The invention describes an isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a specific clone where (I) binds to and neutralises botulinum neurotoxin type A (BoNT/A). An isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a clone chosen from clone 25, C25, C39, IC6, 3D12, B4, IF3, huC25, Az1, Az2, WR1(V), WR1(T), 3-1, 3-8, 3-10 and ING1, where (I) binds to and neutralizes botulinum neurotoxin type A (BoNT/A). Also described are: a polypeptide (II) comprising BoNT/A neutralising epitope having an epitope that is

specifically bound by an antibody expressed by clones as mentioned in (I), producing (I); and a composition (III) comprising several antibotulinum neurocoxin antibodies, where each antibody is specific for a different epitope of a botulinum neurocoxin and the combination of

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RESULT 2
ADR36685
ID ADR3
AC ADR3
AC ADR3
AC ADR3
AC ADR3
AC ADR3
AC MOUS
XX MOUS
KW SONT
KW SONT
KW 11gh
XX US2C
XX US2C
XX US2C
XX US2C
XX US2C
XX US2C
XX MUS
PF 01-J
XX (REC
XX WPI
XX WPI
XX WPI
XX WPI
XX WPI
XX WES
PT WSE
PT WSE
PT WSE
CC The
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Best Local Similarity
Matches 109; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                antibodies shows greater toxin neutralisation than the single antibodies in surplus. The following are disclosed: a pharmaceutical composition comprising (I), and a kit comprising (I) (I) is useful for neutralising south antibody and for neutralising a botulinum neurotoxin which involves contacting neurotoxin with (I) in specific for a different epitope of the botulinum neurotoxin and the combination of antibodies shows greater toxin neutralisation than the single antibodies in surplus. (I) is useful for diagnosing the botulism or for treating pathologies associated with botulinum neurotoxin poisoning. (I) exhibits specificity and affinity towards BoNT/A. (I) enables rapid detection or diagnosis of botulism. This is the amino acid sequence of a mouse light chain variable fragment anti-botulinum toxin
                                                                                                                                                                                                                                                                                                                                                                                                                                                               antibacterial; antibody; botulinum neurotoxin type A; BONT/A; BONT/A neutralising epitope; anti-botulinum neurotoxin antibody; toxin neutralisation; botulinum neurotoxin polsoning; mouse; light chain variable region; single chain antibody; scFv.
                                                                                                     New isolated antibody that neutralizes botulinum neurotoxin type A, useful for diagnosing botulism or for treating pathologies associated
                                                                                                                                                                                              Marks JD,
                                                                                                                                                                                                                                                                    31-AUG-1998; 98US-00144886.
01-AUG-2002; 2002US-0400721P.
                                                 Example 1; SEQ ID NO 87; 110pp; English
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                                                                                                                                                             2004-652009/63.
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                                                                                                                                                                                                Ameradorfer P,
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                                                                                      neurotoxin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Score 559; DB 8
100.0%; Pred. No. 1e-34;
                                                                                      poisoning.
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                                                                                                                                                                                                                                                                                                                                                                                                                     CC clone where (I) binds to and neutralises botulinum neurotoxin type A (CC (BONT/A). An isolated antibody (I) that specifically binds to an epitope CC specifically bound by an antibody expressed by a clone chosen from clone CS5, C25, C39, 1C6, 3D12, B4, 1F5, huC25, Ar1, Ar2, WR1(V), WR1(T), 3-1, C3-8, 3-10 and ING1, where (I) binds to and neutralizes botulinum cc neurotoxin type A (BONT/A). Also described are: a polypeptide (II) comprising BONT/A neutralising epitope having an epitope that is specifically bound by an antibody expressed by clones as mentioned in (I) producing (I), and a composition (III) comprising several anti-cc different epitope of a botulinum neurotoxin and the combination of antibodies hows greater toxin neutralisation than the single antibodies in surplus. The following are disclosed: a pharmaceutical composition (C comprising (I); and a kit comprising (I) (I) is useful for neutralising C BONT/A antibody and for neutralising a botulinum neurotoxin which (I) is specific for a different epitope of the botulinum neutrotoxin and the combination of antibodies shows greater toxin neutralisation than the single antibodies in surplus. (I) is useful for diagnosing the botulism (C specific for a different epitope of the botulinum neutrotoxin than the combination of antibodies associated with botulinum neutrotoxin (I) exhibits specificity and affinity towards BONT/A. (I) cenables rapid detection or diagnosis of botulism. This is the amino acid sequence of a mouse light chain variable fragment anti-botulinum toxin
                                                                                                                                                                                                                                                                                                                   Query Match
Best Local S
Matches 103
                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 107
ADR38673 standard; peptide; 107
                                                                                                                                                                                                                                                                                                                     103;
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                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                    VRFSGSGSGTSYSLTISRMEAEDAATTYYCQQWSSNPLTFGSGTKLELKR
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                                                                                                                        VRFSGSGSGTSYSLTISRMEAEDAATYYCQQWSSNPLTFGSGTKLELKR
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Pred. No. 1.2e-31;
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ALESULT 3
ADR338673
ID ADR3 8673
AC ADR3
AC AD
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01-AUG-2002; 2002US-0400721P.
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The invention describes an isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a specific

New isolated antibody that neutralizes botulinum neurotoxin type

WPI; 2004-652009/63.

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ADW8612
ADW8612
ADW8612
AC ADW8
XX ADW8
XX ADW8
XX Oyte
CAN CYTE
XX OYTE
XX OYTE
XX OYTE
XX ONUS
XX OYTE
XX ADW8
XX OYTE
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Best Local &
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 107
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                                                                                                                                                                                                                                                                                                                                                                                                           apoptosis stimulation; cancer; ovary tumor; uterine cervix tumor; endometroid carcinoma; breast tumor; pancreas tumor; CA6 glycotope; autoimmune disease; graft rejection; viral infection; D86;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADW86126 standard; protein; 107
                                    21-JUL-2003; 2003US-0488447P
                                                                                                         21-JUL-2004; 2004WO-US023340.
                                                                                                                                                                                                                                                                                                                 Mus musculus.
                                                                                                                                                                                                                                                                                                                                                                                  autoimmune
antibody er
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cytotoxin; immunoconjugate; cytostatic; immunosuppressive; virucide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel cytotoxic conjugate-related mouse ayl protein region SeqID46.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21-APR-2005
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               VRFSGSGSGTSYSLTISRMEAEDAATYYCQQWSSNPLTFGSGTKLELKR 109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DIBLTQSPAIMSASPGEKVIMTCSASSSV--SYMYMYQQKPGSSPRLLIYDTSNLASQVP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             88.0%;
llarity 90.8%;
Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 492; DB 8;
Pred. No. 1.1e-29;
2; Mismatches 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
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antiinflammatory; antibacterial; virucide; cytostatic; antipsoriatic; antidiabetic; vasotropic; vaccine; protein purification; protein foldin diagnosis; inflammation; immune disorder; allergic hypersensitivity; infection; autoimmune disease; asthma; georiasis; insulin despendent diabetes; multiple sclerosis; rheumatoid arthritis; systemic lupus erythematosus; myasthenia gravis; hematological disease; neoplasm; immunoglobulin; variable region.

folding;

58

10-FEB-2005

(first entry

Immunoglobulin variable domain protein seqid 54.

14-MAY-2004; 2004WO-GB002102.

WO2004101790-A1

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ADUBGES 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 12; SEQ ID NO 46; 179pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      61 VRFSGSGSGTSYSLTISRMEAEDAATYYCQOWSSNPLTFGSGTKLELKR 109
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DIBLIQSPAIMSASPGBKVTTTCRASSSVS9SYLGWYQQKPGSSPRLLIYDTSNLASGVP 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DIQMTQSPAIMSASPGEKVTMTCSASSSV-JSYMYMYQQKPGSSPRLLIYDSTNLASGVP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Tavares DJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               87.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 491; DB 9;
Pred. No. 1.3e-29;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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                                                                             antiinflammatory; antibacterial; virucide; cytostatic; antipsoriatic; antidiabetic; vasotropic; vaccine; protein purification; protein folding; diagnosis; inflammation; immune disorder; allergic hypersensitivity;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Recovering a polypeptide that unfolds reversibly from a repertoire of polypeptides for treating e.g., cancer, by unfolding a portion of the displayed polypeptides and refolding a portion of the unfolded
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17-MAR-2004; 2004US-0554021P.
infection; autoimmune disease; asthma; psoriasis; insulin dependent diabetes; multiple sclerosis; rheumatoid arthritis; systemic lupus erythematosus; myasthenia gravis; hematological diseas neoplasm; immunoglobulin; variable region.
                                                                                                                                                                Immunoglobulin variable domain protein seqid 60.
                                                                                                                                                                                                        10-FEB-2005
                                                                                                                                                                                                                                                                                   ADU86571 standard; protein; 210
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      87.8%;
88.1%;
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Pred. No. 2.4e~29;
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cc reversibly from a repertoire of polypeptides that unfolds reversibly and common selectable characteristic that distinguishes folded college polypeptides from unfolded or misfolded polypeptides. The method cc comprises: providing a polypeptide display system comprising the crepertoire of displayed polypeptides; unfolding at least a portion of the unfolded cc polypeptides; and recovering at least a portion of the unfolded cc polypeptides; and recovering at least a portion of the unfolded cc portion. The method is useful in recovering a polypeptide that unfolds creversibly and has the selectable characteristic from the refolded cc portion. The method is useful in recovering a polypeptide that unfolds creversibly from a repertoire of polypeptides that unfolds reversibly. The clibrary or repertoire is useful for selecting a polypeptide that crefolded cc polypeptide that unfolds reversibly and comprising an antibody format. The polypeptide that cc antibody variable domain that unfolds reversibly or a polypeptide that cc useful in the manufacture of a medicanent for diagnosing, treating or cytokine receptor, enzyme, enzyme co-factor or DNA binding protein, such case an inflammatory state, allergic hypersensitivity, cancer, bacterial or cytokine receptor, enzyme, enzyme co-factor or DNA binding protein, such case of unitable selectories, rheumatoid arthritis, systemic lugus crythematosus, Crohn's disease, myasthenia gravis, leukemia or solid cumor. This is the amino acid sequence of an immunoglobulin variable in the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Recovering a polypeptide that unfolds reversibly from a repertoire polypeptides for treating e.g., cancer, by unfolding a portion of tdisplayed polypeptides and refolding a portion of the unfolded
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17-MAR-2004; 2004US-0554021P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention describes a method of recovering a polypeptide that reversibly from a repertoire of polypeptides that unfolds reversil
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                                          region associated with the
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210
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                                          protein purification method of the invention.
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                                                                                                    Query Match
Best Local S
Matches 96
 59
                        61
                                                                                                                  Similarity
                                                                   DIBLTQSPAINSASPGEKVTTTCRASSSVSS8YLGWYQQKPGSSPRLLIYDTSNLASGVP
VRFSGSGSGTSYSLTISRMEAEDAATYYCQQWSTYPLTFGAGTKLELKR
                 VRFSGSGSGTSYSLTISRMEAEDAATYYCQQWSSNPLTFGSGTKLELKR
                                                  DIQMTQSPAIMSASPGEKVTMTCSASSSV--SYMYMYQQKPGSSPRLLIYDSTNLASGVP
                                                                                                                87.8%;
                                                                                                    Score 491; DB
Pred. No. 2;4e
7; Mismatches
                                                                                                                DB 8;
                                                                                                                             8; Length 210,
  107
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Sequence

AAR21305
ID AAR2
XX
AC AAR2
XX
DT 21-A
XX
DB Muri

AAR21305 standard;

protein;

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21-MAY-1992 AAR21305;

(first entry)

Murine VL kappa group IV/VI chain t specific for

Fd; bacteriophage; gene III; filamentous; phagemid; capsid; coat; g3p; binding; adsorption; gene VIII; diverse repertoire; specific binding pairs; replicable genetic display package.

pilus;

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Synthetic

Binding-site

Location/Qualifiers

/label

Binding-site Binding-site

90. .98 /label=

CDR3

/label

l= CDR2 CDR1

note=

D-X-G-X-X motif "

10-JUL-1990; 23-JAN-1992 WO9201047-A

90GB-00015198

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                                                                                                               CThe VK sequence is one of 23 (AAR21286-308) expressed from a single chain CC Fv library. The library produces a diverse repertoire of antibody CC fragments specific for 2-phenyl-5-oxazolone (phox). It was prepd. using CC CDNA generated from mRNA from mice immunised with phox coupled to chicked Serum albumin. The VH and VL kappa sequences were separately amplified by CC PCR (AAQ23474-84) and ligated into fdCAT2 (AAQ23463) for expression on CC the phage surface as fusions with gene III. The resulting library of CC vH genes (A-H, AAR21264-71) in a variety of pairings with 7 Vk genes (a-g AR21286-92). Most clones were VH-B combinations so a further CC interarchical library was prepd. by "crossing" VH-B with the Vk clones sequenced. 14 new partners (AAR21293-308) for VH-B were CC identified. Nearly all the Vk genes were "ox-like", only f, (from the CC original library) and h, p, q, and r (from the hierarchical library) were CC values found. This suggests that phage bearing scFv fragments having weak CC values found. This suggests that phage bearing scFv fragments having weak CC culd also be isolated on a basis of antigen affinity. See also AAR21260-307, 309-311; AAR22450, 565-581
                                        Query Match
Best Local S
Matches 97
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   10-JUL-1990;
19-OCT-1990;
12-NOV-1990;
06-MAR-1991;
15-MAY-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mccafferty J, Pope AF
Jackson RH, Holliger
                                                                                                                                                                                                                                                                                                                                                                                                                              Example 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Producing members of specific binding pairs - by expression recombinant host cells with a secreting replicable genetic of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1992-056862/07
                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (MEDI-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CAMB-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CAMBRIDGE ANTIBODY TECHNOLOGY
                                                      Similarity
DIELTQSPAIMSASPGEKVTTTCKASSSVSSSYLGWYQQKPGSSPRLLIYDTSNLASGVP
                                                                                           110 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                            Fig
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             90GB-00015198.
90GB-00022845.
90GB-00024503.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pope AR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     91GB-00010549
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  91GB-00004744.
                                                                                                                                                                                                                                                                                                                                                                                                                              24; 209pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNCIL.
                                                   87.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ΚP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Johnson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Marks
                                        Score 490; DB
Pred. No. 1.6e
4; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         9, K
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Hoogenboom HRJ,
                                                   DB 2;
                                           8
                                                               Length 110;
                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      genetic display
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Griffiths
                                           0
                                           Gaps
                   6
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complementarity determining region; CDR; rheumatoid arthritis; osteoporosis; bone cancer; metastasis; osteolysis; osteoarthritis; psoriasis; inflammatory bowel disease; multiple sclerosis; diabetes; cytostatic; antiarthritic; antiinflammatory; osteopathic; antidiabetic;
The present sequence is that of the light chain variable region of claimed anti-human RANK ligand (RANK-L) murine monoclonal antibody (mAb) 19H22 (isotype 1gG2b/Kappa). The mAb is produced by hybridoma cells obtained by immunising CB6 f1 mice with human RANK-L protein, and fusing spleen cells with X63-Ag8.653 myeloma cells. 19H22 mAb is specific for human RANK-L, having a binding affinity characterised by a dissociation constant of about 10 power -10 M. RANK-L is a member of the tumour necrosis factor family of proteins and a key regulator of the immune system, bone development and homeostasis. The invention provides altered, chimeric and humanised 19H22 antibodies, including Fab or F(ab')2 fragments, antibodies comprising the light and heavy chain variable regions of 19H22, and antibodies comprising the complementarity determining regions of 19H22. These are used in methods for the
                                                                                                                                                                                                                                             A mouse
disease
                                                                                                                                                                                                                                                                                                                                                                                                              21-AUG-2000;
07-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        28-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mus musculus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RANK ligand; RANK-L; monoclonal antibody; mAb; 19H22; antibody;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Anti-RANK ligand mAb 19H22 light chain variable region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    10-JUN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABB75616
                                                                                                                                                                                                                Claim 6; Page 7; 45pp; English.
                                                                                                                                                                                                                                                                                                          WPI;
                                                                                                                                                                                                                                                                                                                                      Sweet RW,
                                                                                                                                                                                                                                                                                                                                                                  (SMIK )
                                                                                                                                                                                                                                                                                                                                                                                                                                                            21-AUG-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO200215846-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Regior
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                light chain;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              antirheumatic; immunosuppressive; antipsoriatic; neuroprotective;
                                                                                                                                                                                                                                                                                                          2002-242022/29.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  μ
                                                                                                                                                                                                                                           monoclonal antibody 19H22 for treating or preventing including rheumatoid arthritis and osteporosis.
                                                                                                                                                                                                                                                                                                                                                                    SMITHKLINE
                                                                                                                                                                                                                                                                                            ABL53530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         VRFSGSGSGTSYSLTISRMEAEDAATYYCQQWSSNPLTFGSGTKLELKR 109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DIBLTQSPAIMSASPGEKVTMTCRASSSVS9SYLNWYQQKSGASPKLWIYSTSNLASGVP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ARPSGSGSGTSYSLTISRMEAEDAATYYCQQRSSYPLTFGAGTKLEIKR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                                              2000US-0226524P.
2000US-0230639P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            2001WO-US026161
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  VL; mouse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              note= "complementarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    'note= "complementarity determining region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          note= "complementarity determining region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                label= CDR2
                                                                                                                                                                                                                                                                                                                                                                    BEECHAM
BEECHAM
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                                                                                                                                                                                                                                                                                                                                        Wattam
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                determining region
                                                                                                                                                                                                                                                               osteopenic
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RESULT 9
ADR38670
ID ADR3
AC A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  र्
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          5
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Best Local S
Matches 98
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         diagnosis, treatment and prevention of osteopenic diseases, including rheumatoid arthritis, osteoporosis, metastatic and primary bone cancer, wear debris induced osteolysis or osteoarthritis, and immune diseases including psoriasis, insulin dependent diabetes, inflammatory bowel disease or multiple sclerosis. Methods are also provided for the recombinant production of the antibodies, using isolated nucleic acids and transfected host (preferably mammalian) host cells, and a method of diagnosing conditions associated with Thi T-cell activity or osteoclast development and activation, in particular those listed above
The invention describes an isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a specific clone where (I) binds to and neutralises botulinum neurotoxin type A (BoNT/A). An isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a clone chosen from clone $25, C25, C39, IC6, 3D12, B4, 1F3, huC25, Ar1, Ar2, WR1(V), WR1(T), 3-1, 3-8, 3-10 and ING1, where (I) binds to and neutralises botulinum neurotoxin type A (BoNT/A). Also described are: a polypeptide (II) comprising BoNT/A neutralising epitope having an epitope that is specifically bound by an antibody expressed by clones as mentioned in (I)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New isolated antibody that neutralizes botulinum neurotoxin type A, useful for diagnosing botulism or for treating pathologies associated with botulinum neurotoxin poisoning.
                                                                                                                                                                                                                                                                                                                                                                                                    Example 1; SEQ ID NO 72; 110pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Marks
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               31-AUG-1998; 98US-00144886.
01-AUG-2002; 2002US-0400721P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-AUG-2003; 2003US-00632706.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mus sp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 antibacterial; antibody; botulinum neurotoxin type A; BONT/A; BONT/A neutralising epitope; anti-botulinum neurotoxin antibody; toxin neutralisation; botulinum neurotoxin poisoning; mouse; toxin neutralisation; botulinum neurotoxin poisoning; mouse; light chain variable region; single chain antibody; scFv.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 129 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (REGC ) UNIV CALIFORNIA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               09-SEP-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mouse light chain variable region scFv seqid 72.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           02-DEC-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2004-652009/63.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2 IELTQSPAIMSASPGEKVTTTCRASSSVSSSYLGWYQQKPGSSPRLLIYDTSNLASGVPV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          98; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RFSGSGSGTSYSLTISRMEAEDAATYYCQQWSSNPLTFGSGTKLELKR 109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               IVLTQSPAIMSASPGEKVTMTCSASSSV--SYMYMYQQKPGSSPRLLIYDTSNLASGVPV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RFSGSGSGTSYSLTISRMEAEDAATYYCQQWSNFPLTFGAGTKLELKR 107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Amersdorfer P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
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Pred. No. 2.2e-29;
3; Mismatches 5,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
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RESULT 10
AAR79867
ID AAR79
XX AAR79
XX AAR79
XX AAR79
XX Singl
CX Singl
CX Humou
XM Phage
XX Humou
XX Clain
XX Clain
XX Humou
XX Clain
XX Clain
XX Humou
XX Clain
XX Clain
XX Clain
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Best Local S
Matches 97
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         botulinum neurotoxin antibodies, where each antibody is specific for a different epitope of a botulinum neurotoxin antibodies in surplus. The following are disclosed; a phramaceutical composition of antibodies shows greater toxin neutralisation than the single antibodies in surplus. The following are disclosed; a pharmaceutical composition comprising (I); and a kit comprising (I). (I) is useful for neutralising a botulinum neurotoxin which is involves contacting neurotoxin with (I) in surplus, where each of (I) is specific for a different epitope of the botulinum neurotoxin and the combination of antibodies shows greater toxin neutralisation than the single antibodies in surplus. (I) is useful for diagnosing the botulism or for treating pathologies associated with botulinum neurotoxin pathologies associated with botulinum neurotoxin poisoning. (I) exhibits specificity and affinity towards BONT/A. (I) enables rapid detection or diagnosis of botulism. This is the amino acid sequence of a mouse light chain variable fragment anti-botulinum toxin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Single chain antibody; antibody; epidermål growth factor receptor; EGFR; tumour; cancer; glioma; melanoma; carcinòma; diagnosis; assessment; phage antibody library.
                                                                                                       Anti-EGFR antibodies and single chain Fv antibody fragments - obtained from phage-antibody libraries, useful for diagnosis and therapy of
                                                                                                                                                                WPI; 1995-336972/43.
N-PSDB; AAT04020.
                                                                                                                                                                                                                                                                                                              17-MAR-1994;
02-DEC-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mus musculus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Anti-EGFR single chain antibody (Clone 10 D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 107
                                                                                                                                                                                                                     Kettleborough AC,
Mitjans F, Rosell
                                                                                                                                                                                                                                                                                                                                                                                                       21-SEP-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                           WO9525167-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         02-JUL-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAR79867;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAR79867 standard; protein; 244
                                                                                                                                                                                                                                                                          (MERE ) MERCK PATENT GMBH
                                                                                                                                                                                                                                                                                                                                                                   16-MAR-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           59
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 DIELTQSPAIMSASPGEKVTTTCRASSSVSSSYLGWYQQKPGSSPRLLIYDTSNLASGVP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DIELTQSPAIMSASPGEKVIMTCSASSSVSHMY -- WYQQKPGSSPRLLIYDTSNLASQVP
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                                                                                                                                                                                                                                                                                                              94EP-00104160.
94EP-00118970.
                                                                                                                                                                                                                                                                                                                                                                   95WO-EP000978
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    86.9%;
                                                                                                                                                                                                                     Bendig MM,
E, Blasco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 486; DB 8;
Pred. No. 3.1e-29;
2; Mismatches 8
                                                                                                                                                                                                                     Ansell KH, F, Piulats
                                                                                                                                                                                                                                        Guessow
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       8; Length 107;
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                                                                                                                                                                                                                                          Adan J;
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Anti-epidermal growth factor receptor antibodies constructed from anti-EGFR Claim 4; Page 60-61; 93pp; English.

(EGFR) single chain antibodies antibody fragments can be used

diagnosis of tumours and assessment of tumour growth in vitro and in vivo. They may also be used in a pharmaceutical compostion for the therapy of e.g. melanomas, gliomas and carcinomas. The antibodies and fragments are derived from mice but are humanised so as to cause minimum reaction against them. They are produced using the phage antibody library. They are produced using the phage antibody AAT04011-T04026 and AAR79858-R79873)

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RRSULT 11
ADT74993
ID ADT74993
AC ADT74
XX ADT74
XX ADT74
XX ADT74
XX AUT01
KW diabe
KW vacci
XX WO200
PD 28-OC
XX WO200
PD 28-FE
XX O4-MP
PR 15-DE
PR 15-DE
PR 28-FE
XX ALLEY
XX HOLEY
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 244 AA;
                                                                                     This invention relates to the novel treatment of an autoimmune disease that utilises an antibody/ autoantigen construct. Specifically, it refers to a peptide autoantigen that is linked to an antibody raised against an antigen presenting cell (APC) receptor. The present invention describes a method for treating diabetes mellitus by inducing an immune tolerance to an autoantigen, which can be chosen from glutamic acid decarboxylase (GAD), insulin or a heat shock protein (HSP), as well as epitopes derived from each thereof or beta cell antigens that are linked to an APC
                                                                                                                                                                                                                                                                                                                                                                                                                                        Treating an autoimmune disease comprises providing an antibody/autoantigen construct containing an autoantigen linked to an antibody to a receptor of an antigen presenting cell and administering the construct to a subject.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 04-MAR-2003; 2003US-0451816P.
15-DEC-2003; 2003US-0529500P.
28-FEB-2004; 2004US-0548385P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               diabetes mellitus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADT74993 standard; protein; 108 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2004-758274/74.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           04-MAR-2004; 2004WO-US006570.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO2004091543-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                autoimmune disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Light chain murine anti-hL-SIGN scFV antibody protein A2LC Seq
receptor antibody. In particular, the antibody recognises a receptor chosen from DC-SIGNR, MHC, toll receptor or mannose receptor amongst others, which is expressed on the surface of the APC. The antibody/
                                                                                                                                                                                                                                                                                                                                                                                Example 9; Fig 8; 71pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (ALEX-) ALEXION PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             61
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DIELTQSPAIMSASPGEKVTTTCRASSSVSSSYLGWYQQKPGSSPRLLIYDTSNLASGVP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      VRFSGSGSGTSYSLTISRMEAEDAATYYCQQWSSNP--LTFGSGTKLELK 108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     hL-SIGN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Kretz-Rommel A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; antibody; antigen presenting cell; APC; immune tolerance; antidiabetic; immunosuppressive;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      86.2%;
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Pred. No. 1.3e-28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Dakappagari N;
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                              autoantigen construct then works to interfere with the interaction between the APC and immune cells such ad autoreactive T cells, and hence inhibits T-cell proliferation. Accordingly, such compositions exhibit antidiabetic and immunosuppressive activities and can be used in the development of vaccines. Note that the single chain antibodies (scFVs) used in this specification contain both variable light and heavy chain regions connected by a linker, their short length makes these antibody fragments particularly suitable for antigen linkage, and the capacity for binding to the receptor is preserved. This polypeptide sequence is a light chain murine anti-human L-SIGN (hL-SIGN) antibody protein of the
                                                                                                                                                                                                                                                                                                                                                                                            Sequence 108
02-DEC-2004
                                     ADR38691;
                                                                        ADR38691 standard; peptide;
                                                                                                                                                                   62
                                                                                                                                                                                                    64
                                                                                                                                                                                                                                                                 4 LTQSPAIMSASPGEKVTTTCRASSSVSSSYLGWYQQKPGSSPRLLIYDTSNLASGVPVRF 63
                                                                                                                                                                                                                                                                                                                   96; Conservative
                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                 SGSGSGTSYSLTISRMEAEDAATYYCQQWSSYPPTFGGGTKLBIKR
                                                                                                                                                                                                SGSGSGTSYSLTISRMEAEDAATYYCQQWSSNPLTFGSGTKLELKR
                                                                                                                                                                                                                                           LTQSPAIMSASPGEKVTMTCSASSSV--SYMYMYQQKPGSSPRLLIYDTSNLASGVPVRF
                                                                                                                                                                                                                                                                                                                                                                                             A,
(first entry)
                                                                                                                                                                                                                                                                                                                                       90.6%;
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در
                                                                                                                                                                                                                                                                                                                   Score 481; DB 8;
Pred. No. 7.4e-29;
2; Mismatches 6
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                                                                                                                                                                                                                                                                                                                                                          Length 108
                                                                                                                                                                                                                                                                                                                       Indels
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                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                   61
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RESULT 12
ADD33691
ID ADD336
XX ADD336
AC ADD3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      31-AUG-1998;
01-AUG-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-AUG-2003; 2003US-00632706
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BONT/A neutralising epitope; anti-botulinum neurotoxin antibody; toxin neutralisation; botulinum neurotoxin poisoning; mouse;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   antibacterial;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (REGC ) UNIV CALIFORNIA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      light chain variable region; single chain antibody; scFv.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       chain variable region scFv seqid 93.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      98US-00144886.
2002US-0400721P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               antibody; botulinum neurotoxin type A; BoNT/A;
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The invention describes an isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a specific clone where (I) binds to and neutralises botulinum neurotoxin type A (BoNT/A). An isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a clone chosen from clone \$25, C25, C39, 1C6, 3D12, B4, 1F3, huC25, Ar1, Ar2, WR1(Y), WR1(T), 3-1, 3-8, 3-10 and ING1, where (I) binds to and neutralizes botulinum neurotoxin type A (BoNT/A). Also described are: a polypeptide (II) New isolated antibody that neutralizes botulinum neurotoxin type A, useful for diagnosing botulism or for treating pathologies associated with botulinum neurotoxin poisoning. Example 1; SEQ ID NO 93; 110pp; English

WPI; 2004-652009/63.

Marks JD,

Amersdorfer

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RESULT 13
AAR79870
ID AAR79
XX AAR79
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XX AAR79
XX Sini
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XX Sini
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XX AAR11
XX AA
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                        Kettleborough AC,
Mitjans F, Rosell
                                                                                                                                         Anti-EGFR antibodies and single chain from phage-antibody libraries, useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      17-MAR-1994;
02-DEC-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           16-MAR-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Anti-EGFR single chain antibody
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      02-JUL-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAR79870 standard; protein;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (MERE )
                                                                                                                                                                                                                                                                                          1995-336972/43.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 DIELTOSPAIMSASPGEKVTTTCRASSSVSSSYLGWYQQKPGSSPRLLIYDTSNLASGVP
                                                                                                                                                                                                                                                      AAT04023.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MERCK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       VRFSGSGSTSYSLTISRMEAEDAATYYCQQWSSNPLTFGSGTKLELKR 109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       VRFSGSGSGTSYSLTISSMEAEDAATYYCQQWSSYPLTFGAGTKVELRR 109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DIELTQSPAIMSASPGEKVTTTCSVSSSISSSNLHWYQQKSGTSPKLWIYGTSNLASGVP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PATENT GMBH.
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94EP-00118970.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           95WO-EP000978
                                                                                                                                                                                                                                                                                                                                                    Bendig MM,
E, Blasco
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Pred. No. 7.5e-29;
                                                                                                                                                                                                                                                                                                                                                        '1
                                                                                                                                                                                                                                                                                                                                                        Ansell KH, F, Piulats
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (Clone 5 F 1).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        carcinoma; diagnosis; assessment;
                                                                                                                                         Fv antibody fragments - obtained for diagnosis and therapy of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 8, Length 109;
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Matches
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Anti-epidermal growth factor receptor (EGFR) single chain antibodies and antibodies constructed from anti-EGFR antibody fragments can be used for diagnosis of tumours and assessment of tumour growth in vitro and in vivo. They may also be used in a pharmaceutical composition for the therapy of e.g. melanomas, gliomas and carcinomas. The antibodies and fragments are derived from mice but are humanised so as to cause minimum reaction against them. They are produced using the phage antibody library. They are produced using the phage antibody 1.34704011-T04026 and AAR79858-R79873)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Single chain antibody; antibody; epidermal growth factor receptor; EGFR; tumour; cancer; glioma; melanoma; carcinoma; diagnosis; assessment; phage antibody library.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Anti-epidermal growth factor receptor (BBFR) single chain antibodies and antibodies constructed from anti-EGFR antibody fragments can be used for diagnosis of tumours and assessment of tumour growth in vitro and in vivo. They may also be used in a pharmaceutical composition for the therapy of e.g. melanomas, gliomas and carcinomas. The antibodies and fragments are derived from mice but are humanised so as to cause minimum reaction against them. They are produced using the phage antibody library. They are produced using the phage antibody labrary. (See AAT04011-T04026 and AAR79858-R79873)
                                                                                                                                                                                                                                                                                                         Claim 4; Page 70; 93pp; English
                                                                                                                                                                                                                                                                                                                                                                                              Anti-EGFR antibodies and single chain Fv antibody fragments - obtained from phage-antibody libraries, useful for diagnosis and therapy of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1995-336972/43.
N-PSDB; AAT04024.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                17-MAR-1994;
02-DEC-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       21-SEP-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mus musculus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Kettleborough AC,
Mitjans F, Rosell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              16-MAR-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO9525167-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Anti-EGFR single chain antibody (Clone 7 G 1).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAR79871;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAR79871 standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (MERE ) MERCK PATENT GMBH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         195 VRFSGSGSGTSYSLTISRMEAEDAATYYCQQNSSYPLTFGAGTKLEIK 242
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             137 DIELTQSPTIMSASPGEKVTMTCSDSSSVSYTY--WYQQKTGSSPRLLIYDTSNLASGVP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            61 VRFSGSGSGTSYSLTISRMEAEDAATYYCQQWSSNPLTFGSGTKLELK 108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 DIELTOSPAIMSASPGEKVTTTCRASSSVSSSYLGWYQQKPGSSPRLLIYDTSNLASGVP
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94EP-00118970
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E, Blasco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  entry)
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Pred. No. 1.6e-28;
3; Mismatches 7
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, Piulats
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19-OCT-1990;
12-NOV-1990;
06-MAR-1991;
15-MAY-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      rd; bacteriophage; gene III; filamentous; phagemid; capsid; coat; g3p; binding; adsorption; gene VIII; diverse repertoire; specific binding pairs; replicable genetic display package.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Binding-site
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The VK sequence is one of 23 (AAR21286-308) expressed from a single chain Fv library. The library produces a diverse repertoire of antibody fragments specific for 2-phenyl-5-oxazolone (phox). It was prepd. using cDNA generated from mRNA from mice immunised with phox coupled to chicked serum albumin. The VH and VL kappa sequences were separately amplified by PCR (AAQ23474-84) and ligated into fdCAT2 (AAQ23463) for expression on
                                                                                                                                                                                                                                                                                                                                                                                                        Mccafferty J, Pope AR, i Jackson RH, Holliger KP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     23-JAN-1992
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                                                                                                                                                                                                                                                                     Producing members of specific binding pairs - by expression in recombinant host cells with a secreting replicable genetic display
                                                                                                                                                                                                                                                                                                                                                   WPI; 1992-056862/07.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (CAMB-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                10-JUL-1990;
                                                                                                                                                                                       Example 22; Fig 24; 209pp; English.
                                                                                                                                                                                                                                               package.
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l Similarity 88.9%;
96; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 90GB-00015198.
90GB-00022845.
90GB-00024503.
91GB-00004744.
91GB-00010549.
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/label= CDR1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       label= CDR2
                                                                                                                                                                                                                                                                                                                                                                                                           Johnson KS,
, Marks JD;
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Pred. No. 1:6e-28;
3; Mismatches 7
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cc the phage surface as fusions with gene II. The resulting library of clones was diverse; 23 hapten binding clones were sequenced revealing 8 cc (H genes (A.H; ARR21264-71) in a variety of pairings with 7 Vk genes (a-g), ARR21286-92). Most clones were VH-B combinations so a further chierarchical library was prepd. by "crossing" VH-B with the Vk repertoire. The resulting library was screened for hapten binding and 24 clones sequenced. 14 new partners (AAR21293-308) for VH-B were consignal library) and h, p, q, and r (from the hierarchical library) were voxil type genes. Of the 24 hierarchical clones, only one was of type crossing. The Kd of VH-B Vk-d for phox-GABA was 10 nM, one of the highest caffinit les can be selected with antigen, probably due to the avidity of the multiple antibody heads on the phage. The different combinations cc could also be isolated on a basis of antigen affinity. See also AAR21260-307, 309-311; AAR22450, 565-581
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    61
                           61 VRFSGSGSGTSYSLTISRMEAEDAATYYCOQWSSNPLTFGSGTKLELKR 109
                                                                                                                                                                94;
                                                                                                                                                                                    Similarity
ARFSGSGSGTSYSLTISSVEAEDAATYYCQQYSGYPLTFGAGTKLEIKR
                                                                               DIELTOSPAINSASPGEKVTMTCRASSSVSSSYLHWYQQKSGASPKLWIYSTSNLASGVP
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86.2%;
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# ALIGNMENT'S

Ig kappa chain V region (F3, anti-AFP) - mouse (fragment) C;Species: Mus musculus (house mouse) C;Date: 10-Nov-1997 #sequence\_revision 23-Jan-1998 #text\_(C;Accession: PC4405 R;Deng, J.B.; Han, H.; Su, C.Z.; Chen, C.Q. Chinese Biochem. J. 12, 648-653, 1996 A;Title: Generation of a phage display library of the imma, Reference number: PC4405 A;Molecule type: mRNA
A;Residues: 1-107 cDEN
A;Cross-references: UNIPROT:Q8K1F0; UNIPARC:UPI00001767BB
A;Experimental source: spleen cell
C;Superfamily: immunoglobulin V region; immunoglobulin homology
F;16-89/Domain: immunoglobulin homology <IMMs RESULT 1 PC4405 문 Ś 片 5 A; Accession: PC4405 Matches Query Match Local 59 ARFSGSGSGTSYSLTISRVEAEDAATYYCQQWRDNPPTFGGGTKLEIKR 107 61 VRFSGSGSGTSYSLTISRMEAEDAATYYCQQWSSNPLTFGSGTKLELKR 109 1 DIELTQSPAIMSASPGEKVTTTCRASSSVS\$SYLGWYQQKPGSSPRLLIYDTSNLASGVP 60 90; Similarity Conservative 82.8%; Score 463; DB 2; Pred. No. 3.4e-32; 6; Mismatches 11 23-Jan-1998 #text\_change 09-Jul-2004 of the immunoglobulin repertoire from huma 11; Indels Length 107; 2 Gaps 88

R.;Kofler, R.; Balderas, R.S.; Jöhnson, M.B.; Noonan, D.J.; Duchosal, M.A.; I. J. Clin. Invest. 82, 852-860, 1988
A;Title: Immunoglobulin kappa light chain variable region gene complex organization and in A;Reference number: A94689; MUID:88331394; PMID:3138286
A;Accession: A32513
A;Accession: A32513
A;Molecule type: DNA
A;Residues: 1-130 < KOF>
A;Residues: 1-130 < KOF>
A;Cross-references: UNIPARC:UPI0000114D9C; GB:M20834; NID:9196943; PIDN:AAA38846.1; PID:C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Superfamily: immunoglobulin V region; Immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin homology <IMM> Ig kappa chain precursor V region (MRL22) - mouse C;Species: Mus musculus (house mouse) C;Date: 21-May-1990 #sequence\_revision 31-Dec-1990 #text\_change 21-Jan-2000 C;Accession: A32513 RESULT 2 A32513 Matches Query Match Local Similarity 91; Conservative 82.6%; Score 462; DB 2; I pred. No. 5.1e-32; 5; Mismatches 11; Length 130; Indels 0 Gaps 0

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Ig light chain V region - mouse (fragment)
N,Alternate names: L6 anti-tumor antibody
C;Species: Mus musculus (house mouse)
C;Date: 27-Apr-1993 #sequence revision 18-Nov-1994 #text_change 21-Jan-2000
C;Accession: A42848; S33902
R;Fell, H.P.; Gayle, M.A.; Yelton, D.; Lipsich, L.; Schieven, G.L.; Marken, J. Biol. Chem. 267, 1552-1558, 1992
A;Title: Chimeric L6 anti-tumor antibody. Genomic construction, expression, A;Reference number: A42848; MUID:92348410; PMID:1639794
A;Accession: A42848
A;Status: preliminary
A;Carana tume, DNA
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(;Species: Mus musculus (house mouse)
(;Species: Mus musculus (house mouse)
(;Date: 28-Oct-1996 #sequence_revision 13-Mar-1997 #text_change 20-Jun-2000
(;Accession: $66536
R;Tsiotis, G.; Hasse, W.; Engel, A.; Michel, H.
Eur. J. Biochem. 231, 823-831, 1995
Eur. J. Biochem. 231, 823-831, 1995
                                                                                                                                                                                                                                                A;Molecule type: mRNA
A;Residues: 1-120 <TSI>
A;Cross-references: UNIPARC:UPI00001137B0; EMBL:X88903; NID:g895870;
C;Superfamily: immunoglobulin V region; immunoglobulin homology
F;16-91/Domain: immunoglobulin homology <IMM>
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A;Note: sequence extracted from NCBI backbone (NCBIN:109958, NCBIP:109959)
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: immunoglobulin
F;16-89/Domain: immunoglobulin homology <IMM>
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A; Residues: 1-107 < FEL>
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        ARPSGSGSGTSYSLTIGTMEAEDVATYYCQQGSSIPFTFGGGTKLEIKR
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                                                                                                                                                                      Conservative
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83.3%;
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80.7%;
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                                                                                                                                                                                       Score 455; DB 2; Length 120, Pred. No. 1.8e-31;
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Pred. No. 1.3e-31;
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                                                                                                                                                                                                                                                                                            PH
    Ig kappa chain V regions (27.7.2 and 27.4b.2) C;Species: Mus musculus (house mouse) C;Date: 23-Mar-1989 #sequence_revision 23-Mar-
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RESULT 7
A30562
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S04573
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C/Species: Mus musculus (house mouse)
C/Species: Mus musculus (house mouse)
C/Species: Mus musculus (house mouse)
C/Accession: B32456
C/Accession: B32456
R/Dombrink-Kurtzman, M.A.; Johnson, L.S.; Riordan, G.S.; Bedzyk, W.D.; Voss Jr., E.W.
J. Biol. Chem. 264, 4513-4522, 1989
A;Title: Variable region primary structures of a high affinity anti-fluorescein immunoglo
A;Reference number: A32456; MUID:89174706; PMID:2494173
A;Accession: B32456
A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-130 <DOM>
A;Residues: THIORDAY
A;R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Cross-references: UNIPARC:UPI0000115DFB; EMBL:X14620; NID:952031; PID A;NOte: the authors translated the codon Acf for residue 47 as Asn C;Superfantly: immunoglobulin V region; immunoglobulin homology C;Keywords: heterotetramer; immunoglobulin C;Keywords: heterotetramer; immunoglobulin C;Keywords: heterotetramer; immunoglobulin F;1-22/Domain: signal sequence #status predicted <SIG> F;2-3-130/Product: Ig kappa chain V region (fragment) #status predicted F;38-131/Domain: immunoglobulin homology <IMM>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  R;Kofler, R.; Noonan, D.J.; Strohal, R.; Baldèras, R.S.; Moller, N.P.H.; Dixon, Bur. J. Immunol. 17, 91-95, 1987

Bur. J. Immunol. 18, 91-95, 1987

A;Title: Molecular analysis of the murine lupus-associated anti-self response: 1
A;Reference number: S04573; MUID:87133856; PMID:3102255

A;Accession: S04573.
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C;Date: 04-Dec-1992 #sequence_revision 04-Dec-1992
C;Accession: S04573
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A; Molecule type: mRNA
A; Residues: 1-130 < KOF>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ig kappa chain
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Best Local S
Matches 89
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Best Local S
Matches 89
                                                                                                                                                                                                                                                            62
                                                                                                                                                                                                                                                                                                                                                  24 IVLTQSPAIMSASPGEKVTMTCSASSSVSSKYLNWYQQRSGASPKLWIYGTSNLASGVPA 83
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   64
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     , 68
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  h 81.0%;
Similarity 83.2%;
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                                                                                                                                                                  RFSGSGSGTSYSLTISRMEAEDAATYYCQQWSSNPLTFGSGTKLELK 108
                                                                                                                                                                                                                                                                                                                                                                                                        IELTQSPAIMSASPGEKVTTTCRASSSVSSSYLGWYQQKPGSSPRLLIYDTSNLASGVPV 61
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Pred. No. 2.9e-31;
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Pred. No. 2e-31;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
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23-Mar-1989

#text\_change

21-Jan-2000

mouse

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C;Accession: A30562
R;Sikder, S.K.; Borden, P.; Gruezo, F.; AKOLDAL,
J. Immunol. 142, 888-993, 1989
A;Title: Amino acid substitutions in V-H CDR2 change the idiotype but :
A;Reference number: A30562; MUID:89110066; PMID:2464031
A;Accession: A30562
A;Accession: A30562
A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-107 <SIK>
A;Resi
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A;Cross-references: UNIPARC:UPI0000176780
C;Superfamily: immunoglobulin V region; in
C;Keywords: heterotetramer; immunoglobulir
F;16-89/Domain: immunoglobulin homology <
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Nucleic Acids Res. 12, 4229-4246, 1984
A;Title: Immunoglobulin gene 'remnant' DNA
A;Reference number: S03410; MUID:84221356;
A;Accession: S03410
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ig kappa chain precursor V region (ABE8) - C:Species: Mus musculus (house mouse)
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A;Note: this sequence was determined from the nonfunctional differentiated A;Note: this sequence was determined from the nonfunctional differentiated C;Superfamily: immunoglobulin v region; immunoglobulin homology C;Keywords: heterotetramer; immunoglobulin C;Leywords: heterotetramer; immunoglobulin F;1-6/Domain: signal sequence (fragment) #status predicted <SIG>F;7-113/Product: Ig kappa Chain V region ABES #status predicted <MAT>
                                                                                                     R,Sikder, S.K.; Borden, P.; Gruezo, P.; Akolkar, P.N.; Bhattachary J. Immunol. 142, 888-893, 1989
A;Title: Amino acid substitutions in V-H CDR2 change the idiotype
                                                                                                                                                                                                           Ig kappa chain V region (27.10.2) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 23-Mar-1989 #sequence_revision 23-Mar-1989 #text_change 21-Jan-2000
C;Accession: B30562
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A;Residues: 1-113 <SEL>
                              A;Reference number: A30562; MUID:89110066; PMID:2464031
A;Accession: B30562
A;Status: preliminary
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        61 VRFSGSGSGTSYSLTISRMEAEDAATYYCQQWSSNPLTFGSGTKLELKR 109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ผ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 DIELTQSPAIMSASPGEKVTTTCRASSSVSSSYLGWYQQKPGSSPRLLIYDTSNLASGVP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  91;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RFSGSGSGTSYSLTISRMEAEDAATYYCOOWSSNPLTFGSGTKLELKR 109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RFSGSGSGTSYSLTISSMEAEDAATYYCQQWSSNPYTFGGGTKLEIKR 107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  IELTQSPAIMSASPGEKVTTTCRASSSVSSSYLGWYQQKPGSSPRLLIYDTSNLASGVPV 61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ARFSGSGSGTSYSLTINSMEAEDAATYYCQQWST--LTFGAGTKLELKR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BIVLTQSPAIMSASPGERVTMTCSASSSVSSSYLYWYQQKSGSSPKLWIYSISNLASGVP
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                                                                                                                                                  P.; Gruezo, F.; Akolkar, P.N.; Bhattacharya,
3, 1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         80.9%;
83.5%;
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Pred. No. 2.9e-31;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 452; DB 2
Pred. No. 3e-31;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
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PMID:6328416
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2; Length 113;
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                                                                                                                                                                                        S.B.; Morrison,
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Ig kappa chain precursor V region (anti-phenyloxazolone 6F6) - mouse C;Species: Mus musculus (house mouse) C;Date: 31-Dec-1991 #sequence revision 09-Aug-1996 #text_change 16-Jul-1999 C;Accession: JL0079; A49044; B49044 B49044 R;Kaartinen, M:;Rocca-Serra, J:; Maekelae, O. Mol. Immunol. 25, 859-865, 1988 Mol. Immunol. 25, 859-865, 1988 A;Title: Combinatorial association of V genes: one VH gene codes for three r A;Reference number: JL0076; MUID:89096973; PMID:3211160
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Molecule type: mRNA
A;Residues: 1-107 <SIK>
A;Cross-references: UNIPARC:UPI00001767BA
C;Superfamily: immunoglobulin V region; immuno
C;Keywords: heterotetramer; immunoglobulin
F;16-89/Domain: immunoglobulin homology <IMM>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Cross-references: UNIPARC:UPI000017371F; GB.M27792; NID:g197159
A;Experimental source: mRNA clones for anti-phenyloxazolone antibody 6F6
A;Note: the authors translated the codon TTC for residue 8 as Pro and TTC for residue 10
A;Note: the codon given for 1-Met (AGT) is inconsistent with the authors' translation
R;Milstein, C.; Even, J.; Jarvis, J.M.; González-Fernandez, A.; Gherardi, B.
Eur. J. Immunol. 22, 1627-1634, 1992
A;Title: Non-random features of the repertoire expressed by the members of one V kappa 90
A;Reference number: A49044; MUID:92289826; PMID:1601044
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                                                                                                                                                                                                                     F;1-22/Domain: signal sequence #status predidted <SIG>
F;23-130/Product: Ig kappa chain V region (6%6) #status
F;38-111/Domain: immunoglobulin homology <IVNS
F;45-109/Disulfide bonds: #status predicted
                                                                                                                                                                                                                                                                                                                                 C;Complex: An immunoglobulin heterotetramer subunit consists of two identical light hain disulfide bonds. In some cases, such as IgA and IgM, the subunits associate int C;Superfamily: immunoglobulin V region; immunoglobulin homology C;Keywords: heterotetramer; immunoglobulin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Experimental source: BALB/c germ-line
A;Note: sequences extracted from NCBI backbone; partial nucleotide sequences of 13
A;Note: sequence extracted from NCBI backbone (NCBIN:106807, NCBIP:106822)
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A; Introns: 17/1
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A;Residues: 114-116 <MIL2>
A;Cross-references: UNIPARC;UPI000011E8B5; GB:S37664; NID:g250215; PIDN:AAB22332.1; PID:g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Cross-references: UNIPARC:UPI000011760C; GB:S37663; NID:g250214; PIDN:AAB22331.1;
A;Note: sequence extracted from NCBI backbond (NCBIN:106802, NCBIP:106809)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; Molecule type: DNA
A; Residues: 1-25 < MIL>
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A; Residues: 1-130 < KAA>
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Best Local
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                                                         IELTQSPAINSASPGEKVTTTCKASSSVSSSYLGWYQQKPGSSPRLLIYDTSNLASGVPV 61
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Pred. No. 3.5e-31;
                                                                                                                                                Score 450; DB 1;
Pred. No. 5.1e-31
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                                                                                                                                                                        Length 130
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C;Date: 19-Mar-1997 ##EQUESTORM (C;Date: 19-Mar-1997) ##EQUESTORM (C;Date: 19-Mar-1997) ##EQUESTORM (C;Date: S25058) ##ERSON (C;Date: S25058) ##ER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        submitted to the EMBL Data Library, A;Reference number: S26459
A;Accession: S29591
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ig kappa chain V region - mouse
C;Species: Mus musculus (house mouse)
C;Date: 06-Jan-1995 #sequence_revision
C;Accession: S29591
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Cross-references: UNIPARC:UPI0000116096; EMBL:X67211; NID:g54828; PIDN:CAA47650.1; PIDC;Superfamily: immunoglobulin V region; immunoglobulin homology F;38-111/Domain: immunoglobulin homology <IMM>
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Ç;Accession:
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A; Residues: 1-103 < KAV>
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       g kappa chain precursor V-J region (38Cl3-V4) - mouse (fragment)
;Species: Mus musculus (house mouse)
;Date: 30-Sep-1991 #sequence_revision 30-Sep-1991 #text_change 21-Jan-2000
;Accession: S05269; JL0064; S03847
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Species: Mus musculus (house mouse);Date: 19-Mar-1997 #text_change 21-Jan-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Status: preliminary
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      RFSGSGSGTSYSLTISRMEAEDAATYYCQQWSSNPLTFGSGTKLELKR 109
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               IELTQSPAIMSASPGEKVTTTCRASSSVS8SYLGWYQQKPGS8PRLLIYDTSNLASGVPV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          IVLTQSPAIMSASPGEKVTMTCSASSSVSK--MQWYQQKSGTSPKRWIYDTSKLASGVPG
                                                                                                                                                                                                                                                                                                              RFSGSGSGTSYSLTISSMEAEDAATYYCQQWSSNPLTFGAGTKLE
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84.8%;
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Pred. No. 8.8e-31;
4; Mismatches 10
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A;Status: preliminary
A;Molecule type: mRNA
A;Molecule type: mRNA
A;Residues: 2-104 <KA3>
A;Cross-references: UNIPARC:UPI0000115F55; EMBL:X59092; NID:g52223; PIDN:CAA41818.1;
A;Reference number: A33730; MUID:89367325; PMID:2505260
A;Accession: B33730
A;Molecule type: DNA
A;Residues: 1-96 <LAM>
A;Residues: 1-96 <AAA38913.1;
A;Rote: the authors translated the codon ATT for residue 2 as Ala
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Reywords: heterotetramer; immunoglobulin
C;Reywords: heterotetramer; immunoglobulin
E;16-91/Domain: immunoglobulin homology <IMM>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ig kappa chain V region - mouse
C;Species: Mus musculus (house mouse)
C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 21-Jan-2000
C;Accession: S29581; S26477; S29589; B33730; B26476
R;Kavaler, J.
submitted to the EMBL_Data Library, April 1991
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A;Cross-references: UNIPARC:UPI0000115DED; EMBL:X14099; NID:g52568; PIDN:CAA32261.1;
A;Cross-references: CUBPROCOLLERY, R.; Levy, S.
R;CArroll, W.L.; Starnes, C.O.; Levy, R.; Levy, S.
J. Exp. Med. 168, 1607-1620, 1988
A;Title: Alternative V kappa gene rearrangements in a murine B cell lymphoma. An expl. A;Reference number: UL0061; MUID:89035985; PMID:3141553
A;Accession: JL0064
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A;Residues: 1-121 <CAR>
A;Residues: 1-121 <CAR>
A;Cross-references: UNIPARC:UPI0000176799; EMBL:X14099
C;Superfamily: immunoglobulin V region; immunoglobulin homology
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A,Reference number: 805267
A,Accession: 805269
A,Molecule type: mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Molecule type: mRNA
A; Residues: 1-106 < KA2>
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A; Accession: S26477
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A; Accession: S29581
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; Accession: S29589
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Pred. No. 1e-30;
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Ig light chain V region - mouse
C;Species: Mus musculus (house mouse)
C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 20-Jun-2000
C;Accession: S38720
C;Accession: S38721
A;Reference number: S38713
A;Reference number: S38713
A;Recession: S38720
A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-108 <CIM>A;Residues: 1-108 <CIM>A;Residues: 1-108 <CIM>C;Superfamily: immunoglobulin
C;Superfamily: immunoglobulin
C;Keywords: immunoglobulin homology <IMM>
F;16-89/Domain: immunoglobulin homology <IMM>
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S38720
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Search completed: November 21, 2005, 12:22:13 Job time : 27.5513 secs
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                                                                                                                                                                                                                                         Query Match 79.5%; Score 444.5; DB 2; Length 108; Best Local Similarity 82.7%; Pred. No. 1.2e-30; Matches 91; Conservative 7; Mismatches 9; Indels 3
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                                                                               61 VRFSGSGSGTSYSLTISRMEAEDAATYYCQQWSSN-PLTFGSGTKLELKR 109
59 LRFSGSGSGTSYSLTISSMEAEDAATYYCQQWSSNPPLTFGAGTKLQLKR 108
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Maximum Match 100%
Listing first 45 summaries
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       Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.
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Q569YB MOUSE

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Q8K1P0 MOUSE

Q8VID10 MOUSE

KV4A MOUSE

Q9UL78 MOUSE

Q9UL78 MOUSE

C9UL78 MOUSE

KV4B MOUSE

KV5B HUMAN

KV6B MOUSE

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62 SC	64 SC	4 []	4 2	Match Local Simi	NON TER SEQUENCE		20,0	L; AF207 P; P0167 ; Q9U410	ngguo Ji 257-259(	loning a	Song X.T., Huang H.L.,	LEOTIDE Med=1256	Muridae; Murinae; NCBI_TaxID=10090; [1]	Eukaryota; Mammalia; E		2000	Q9U410; 01-MAY-2000	NOUSE MOUSE MOUSE		367.5 367.5	370.5	371.5	375	379	382.5	
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BLK 106	BLK 108	RLLIYDTSNLASGVPVRF 61	RLLIYDTSNLASGVPVRF 63	ngth 106; Indels 2; Gaps 1;	64;				Za Zhi	variable region Schistosoma	Xiong Y., Yin C.C.,			Buteleostomi; tia; Sciurognathi;	E)	ibody NP30				homo homo	Q9u170 homo sapien Q6pit5 homo sapien	homo	homo	homo		

RESULT 2

QBKIF1 MOUSE PRELIMINARY;

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Created)
Last sequence update)

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PIR; PH1058; PH1058.

H8SP; P01837; 25C8.

SMR; Q8KLF1; 2-114.

Ensembl; ENSMUSG00000059896; Mus musculus.

InterPro; IPR007110; Ig-1ike.

InterPro; IPR007110; Ig-v.

SMART; SM00406; IGv; 1.

PROSITE; PS50835; IG_LIKE; 1.
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01-JUN-2003 (TrEMBLrel. 24, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
01-mAR-2004 (TrEMBLrel. 26, Last annotation update)
Imunoglobulin gamma-3 kappa chain precursor (Fragment).
Mus musculus (Mouse).
Bukaryota; Metazoa; Chordáta; Craniata; Vertebrata; Euteleostomi;
Bukaryota; Metazoa; Chordáta; Craniata; Glires; Rodentia; Sciurognat
                     NUCLECTIDE SEQUENCE.

STRAIN-MRL/MpJ-lpr/lpr; TISSUE-Spleen;
MEDLINE-93156722; PubMed-8429833; DOI=10.1016/0161-5890(93)90089-T;
MEDLINE-93156722; PubMed-8429833; DOI=10.1016/0161-5890(93)90089-T;
Takahashi S., Itoh J., Nose M., Ono M., Yamamoto T., Kyogoku M.;

"Cloning and cDNA sequence analysis of nephritogenic monoclonal antibodies derived from an MRL/lpr lupus mouse.";
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Anti-VIPase light chain variable region (Fragment).
Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;
Muridae; Murinae; Mus.
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STRAIN-BALB/c; TISSUE-Hyperimmunized spleen;
Zhou Y.-X., Taguchi H., Planque S., Karle S.,
Submitted (MAY-2002) to the EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Q811C3 MOUSE PRELIMINARY;
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MEDLINE=92381444; Pu
Tillman D.M., Jou N.
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Proc. Natl. Acad. Sci. U.S.A. 86:4624-4628(1989)
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4; Mismatches 12;
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EMBL, D14629; BAA03482.1; -; mRNA.
HSSP, P01679; ZFBJ.
SMR; Q811C3; 23-131.
Ensembl; ENSMUSG0000058987; Mus mus
InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig-v.
SMART; SM0406; IGV; 1.
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13-AUG-1987 (Rel. 05, Last sequence update)
10-MAY-2005 (Rel. 47, Last annotation update)
19 kappa chain V-VI region NQ2-6.1.
Mus musculus (Mouse).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi; Muroidea; Muridae; Murinae; Mus.
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HSSP; Q91W12; 1AY1.
SMR; P04945; 1-108.
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                                                                                                                                                                                            InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig v.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG LIKE; 1.
Hybridoma; Immunoglobulin dom
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                                                                                                                                                       REGION
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72 MOUSE
72 MOUSE
QRK1F2,
QBK1F2;
QBK1F2;
QCT-2002 (TrEMBLrel. 22, Created)
01-OCT-2002 (TrEMBLrel. 22, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Anti-VIPase light chain variable region (Fragment).
Mus musculus (Mouse).
Rukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;
Muridae; Murinae; Mus.
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"Two murine natural polyreactive autoantibodies are encoded
"Tonmutated germ-line genes.";
Proc. Natl. Acad. Sci. U.S.A. 86:4624-4628(1989).
EMBL, AP516283; AAM64201.1; -; Genomic_DNA.
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STRAIN-BALB/C; TISSUB-Hyperimmunized spleen;
STRAIN-BALB/C; TISSUB-Hyperimmunized spleen;
Zhou Y.-X., Taguchi H., Planque S., Karle S.,
Zhou Y.-X., Taguchi H., Planque S., Karle S.,
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InterPro; IPR0077110; Ig-like.
InterPro; IPR003596; Ig-v.
SMART; SM00406; IG-v; 1.
PROSITE; PS50835; IG_LIKE; 1.
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HSSP; P01837; 25C8.
SMR; Q8K1F2; 2-112.
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PubMed=2499887;
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     Last sequence update)
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           RESULT
Q5XFY8
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RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Alleschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Alleschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Schaefel M.J., Ugdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA Bosak S.A., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Walla N., Sodergren E.J., Lu X., Gibbs R.A.,
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
RA Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
RA Butterfield Y.S.N., Krzywinski M.I., Skalska U., Smailus D.E.,
RA Rodriguez A.C., Grimwood J., Schmutz J.M., Warrs R.M.,
RA Schnerch A., Schein J.E., Jones S.J.M., Warrs M.A.;
The College CDNA sequences.",
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Best Local Similarity
Matches 89; Conserv
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InterPro; IPR007110; IG-1ike.
InterPro; IPR003597; IG_c1.
InterPro; IPR003597; Ig_mtC.
InterPro; IPR003596; Ig_w.
InterPro; IPR003596; Ig_w.
Pfam; PP07554; C1-set; I.
SMART; SM00409; IG; 2.
SMART; SM00400; IGv; 1.
SMART; SM00406; IGv; 1.
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SEQUENCE
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Mammalia; Eutheria; Euarchontoglires; Glires; Rodent
Muroidea; Muridae; Musimae; Mus.
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STRAIN=FVB/N; TISSUE
MEDLINE=22388257; Pu
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Name=Igk-C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Submitted (MAR-2005) to the EMBL/GenBank/DDBJ databases. EMBL; BC091738; AAH91738.1; -; mRNA.
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                                                                                                                      Q5XFY8 MOUSE
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STRAIN=FVB/N; TISSUE
           Hypothetical
                                                                                                                                                                             MOUSE
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GO:0003823; F:antigen binding; IEA.
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PS00290; IG_MHC; UNKNOWN_1.
235 AA; 25719 MW; BE4E4
                                                                                                                                                                                                                                                                                                                                   RFSGSGSGTSYSLTISRMEABDAATYYCQQWSSNPLTFGSGTKLELKR
                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                        (TrEMBLrel.
                                                                                               (TrEMBLrel.
                                                                                                                                                     PRELIMINARY;
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                                        Created)
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Last annotation update)
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Pred. No. 6.7e-36;
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                                                                                                                                                        PRT;
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RA Strausberg R.L., Feingold B.A., Grouse L.H., Derge J.G.,
RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Altschul S.F., Jordan H., Moore T., Max S.I., Wang J., Heich F.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Heich F.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casawant T.L., Scheetz T.E.,
RA Brownstein M.J., Ugdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Brownstein M.J., WcKernan K.J., Malek J.A., Gunaratne P.H.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Villalon D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,
RA Villalon D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Hotesley R.W., Touchman J.W., Green B.D., Dickson M.C.,
RA Butterfield Y.S.N., Krzywinski M.I., Skalska U., Smailus D.E.,
RT "Generation and initial analysis of more than 15,000 full-length human
RT and mouse cDNA sequences.";
RT Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903 (2002).
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InterPro; IPR007110; Ig-11ke.
InterPro; IPR003597; Ig_c1.
InterPro; IPR003597; Ig_MHC.
InterPro; IPR003596; Ig_v.
Pfam; PF07654; C1-set; 1.
KV6F MOUSE
P04940;
13-AUG-1987
13-AUG-1987
10-MAY-2005
Ig kappa cha:
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NCBI_TaxID=10090;
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SEQUENCE 2:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PROSITE; PS50835; IG_LIKE; 2.
PROSITE; PS00290; IG_MHC; UNKNOWN_1.
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SMART; SM00407; IGc1; 1.
SMART; SM00406; IGv; 1.
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SMR; Q5XFY8; 23-235.
GO; GO:0003823; F:antigen binding; IEA.
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Director MGC Project;
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STRAIN=FVB/N; TISSUI
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                                         kappa chain
        musculus
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                                 7 (Rel. 05, Created)
7 (Rel. 05, Last sequence of the control of t
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235 AA; 2
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        (Mouse)
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                                                                                                                                                                                                                                                   STANDARD;
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81.5%;
                                         sequence update) annotation update) NQ2-17.4.1.
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Pred. No. 1.3e-35;
B; Mismatches 10;
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RESULT 9

RY6I MOUSE
ID TV61 MOUSE
AC P04943;
DT 13-AUG-1987
DT 13-AUG-1987
DT 10-MAY-2005
DE 1g kappa cha
OS Mus musculus
OC Elkaryota; h
OC Mammalia; El
OC Muroidea; Ml
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RA Kaartinen M.
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Best Local :
           NUCLEOTIDE SEQUENCE.
MEDLINE=83271467, PubMed=6877353,
Kaartinen M., Griffiths G.M., Markham A.F., Milstein
Manna sequences define an unusually restricted IgG re
phenyloxazolone and its early diversification.";
Nature 304,320-324(1983).
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                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurogna Muroldea; Muridae; Murinae; Mus.
                                                                                                                                                                     13-AUG-1987 (Rel. 05, Created)
13-AUG-1987 (Rel. 05, Last sequence up
10-MAY-2005 (Rel. 47, Last annotation
Ig kappa chain V-VI region NQ6-8.3.1.
Mus musculus (Mouse)
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Ensembl; ENSMUSG00000062047; Mus
Interpro; IPR007110; Ig-11ke.
Interpro; IPR003596; Ig_v.
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MEDILINE=83271467, PubMed=6877353;
MEDILINE=83271467, PubMed=6877353;
Kasrtinen M., Griffiths G.M., Markham A.F., Milstein
"mRNA sequences define an unusually restricted IgG re
phenyloxazolone and its early diversification.";
                                                                                                           NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hybridoma;
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Mammalia; I
  FUNCTION: Anti-2-phenyl oxazolone
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P01679; 2FBJ.
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Pred. No. 8
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Complementarity-determining-1.
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13-AUG-1987 (Rel. 05, Created)
13-AUG-1987 (Rel. 05, Last seq
10-MAY-2005 (Rel. 47, Last ann
19 kappa chain V-VI region NQ5
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InterPro; IPR003596; Ig_v.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 1.
Hybridoma; Immunoglobulin domain; Immunoglobulin V
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HSSP; P01679; 2FBJ.
SMR; P04943; 1-107.
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                                                                                                                                                                                                                                                                                                                                                                                                 NUCLEOTIDE SEQUENCE.

MEDILINE-83371467; Pubmed=6877353;

MEDILINE-83371467; Pubmed=6877353;

MARKham A.F., Milstein

MEDILINE-837, Milstein

MEDILINE-837, Milstein

MEDILINE-837, Milstein

MEDILINE-837, Milstein

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MEDILINE-837, Markham A.F., Milstein

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Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalla; Butheria; Buarchontoglires; Glires; Rodentia; Sciurognathi;
Muroidea; Muridae; Murinae; Mus.
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                                                         EMBL; K00739; AAA38684.1;
HSSP; P01679; 2FBJ.
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Pred. No. 2.5e
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InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig-v.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG-LIKE; 1.
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PIR; PH1085; PH1085.
HSSP; P01837; 25C8.
SMR; Q8K1F3; 2-112.
                                                                                                                                                                                                                                                                                                                                                                                                          PubMed=249900;
Baccala R., Quang T.V., Gilbert Pagaccala R., Guille Pagacca
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRAIN-BALE/c; TISSUE=Hyperimmunized spleen; Zhou Y.-X., Taguchi H., Planque S., Karle S., Submitted (MAY-2002) to the EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurogna Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        01-OCT-2002 (TrEMBLrel. 22, Created)
01-OCT-2002 (TrEMBLrel. 22, Last sequence update)
01-WAR-2004 (TrEMBLrel. 26, Last annotation update)
Anti-VIPase light chain variable region (Fragment).
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SMART; SM00406; IGV;
PROSITE; PS50835; IG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          QBK1F3 MOUSE PRELIMINARY;
                                                                                                                                                                                                                                           MEDLINE=92381444; PubMed=1512540; DOI=10.1084/jem.176.3.761; Tillman D.M., Jou N.T., Hill R.J., Marion T.N.; "Both IgM and IgG anti-DNA antibodies are the products of cleelective B cell stimulation in (NZB x NZW)F1 mice."; J. Bxp. Med. 176:761-779(1992).
J. Bxp. Med. 176:761-779(1992).
BXBL, AF515282; AAM64200.1; -; Genomic_DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mus musculus (Mouse).
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00406; IGv; 1.
PS50835; IG LIKE; 1.
; Immunoglobulin doma
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               domain; Immunoglobulin V region.
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Pred. No. 3.1e-35
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Complementarity-determining-3.
Framework-4.
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13-AUG-1987 (Rel. 05, Last sequence update)
10-MAY-2005 (Rel. 47, Last annotation update)
19 kappa chain V-VI region NQ2-48.2.2.
Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;
Muroidea; Muridae; Murinae; Mus.
NCBI TaxID=10090;
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InterPro; IPR0077110; Ig-1ke.
InterPro; IPR003796; Ig-v.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG-LIKE; 1.
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MEDLINE-83271467; PubMed-6877353;
MEDLINE-83271467; PubMed-6877353;
Kaartinen M., Griffiths G.M., Markham A.F., Milstein
"mRNA sequences define an unusually restricted IgG re
phenyloxazolone and its early diversification.";
Nature 304:320-324(1983)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EMBL; K00737; AAA38682.1;
HSSP; Q91W12; 1AY1.
SMR; P04941; 1-103.
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Complementarity-determining-1.
Framework-2.
Complementarity-determining-2.
Framework-3.
Complementarity-determining-3.
Framework-4.
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Pred. No. 3.3e-35;
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Pred. No. 3.9e-35;
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Q569Y8_MOUSE PRELIMINARY;
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10-MAY-2005 (TrEMBLrel. 30,
10-MAY-2005 (TrEMBLrel. 30,
10-MAY-2005 (TrEMBLrel. 30,
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13-AUG-1987 (Rel. 05, Created)
13-AUG-1987 (Rel. 05, Last sequence update)
10-MAY-2005 (Rel. 47, Last annotation update)
10-MAY-2005 (Rel. 47, Last annotation update)
17 kappa chain V-VI region NQ5-78.2.6.
Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Sciurognathi;
Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=83271467; PubMed=6877353;
Kaartinen M., Griffiths G.M., Markham A.F., Milstein
"mRNA sequences define an unusually restricted IgG re
phenyloxazolone and its early diversification.";
Nature 304,320-324(1983).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This Swiss-Prot entry is copyright. It is produced through a continuous the Swiss Institute of Bioinformatics and the EMBL the European Bioinformatics Institute. There are no restrict
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
SMART; SM00406; IGv; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ensembl; ENSMUSG00000062047; Mus
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; Q91W12; 1AY1.
P04944; 1-106.
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                                                                                                                                                                  RFXGSGSATSYSLTITSMQAEDAATYYCQQW6SNPLTFGSGTKLEXKR
                                                                                                                                                                                               RFSGSGSGTSYSLTISRMEAEDAATYYCQQW&SNPLTFGSGTKLELKR
                                                                                                                                                                                                                                                                 IBLTQ8PAIMSASPGBKVTTTCRASSSYSSSYLGWYQQKPGSSPRLLIYDTSNLASGVPV
                                                                                                                                                                                                                                              S50835; IG LIKE;
Immunoglobulin c
 S (TrEMBLrel. 30, TrEMBLrel. 30, TrEMBLrel. 30,
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   Last
Last
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Pred. No. 3.9e-35;
6; Mismatches 13
                                      Created)
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Complementarity-determining-2.
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 sequence update)
annotation update)
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RESULT 15
QBKLF0 MOUSE
ID QBKLF0 M
AC QBKLF0 TO 10-CCT-2
DT 01-MAR-2
DT 01-MAR-2
DT 01-MAR-2
DT 01-WAR-2
CO Mus musc
OC Mammalis
OC Muridae;
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RP C STRAIN-FYB/N, TISSUB-Kidney;

RX MEDLINE-22388257, Pubmed=12477932; DOI=10.1073/pnas.242603899;

RX MEDLINE-22388257, Pubmed=12477932; DOI=10.1073/pnas.242603899;

RX Klausner R.D., Collins F.S., Wagner L., Shemmen C.M., Schuler G.D.,

RA Klausner R.D., Collins F.S., Wagner L., Shemmen C.M., Schuler G.D.,

RA Altechul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

RA Villalon D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,

RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

RA Rahey J., Helton B., Ketteman M., Madan A., Rodrigues S., Sanchez A.,

RA Rahersley R.W., Touchman J.W., Green E.D., Dickson M.C.,

RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,

RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,

"Generation and initial analysis of more than 15,000 full-length human

RT and mouse cDNA sequences ";

Lycc. Natl. Acad. Sci. U.S.A. 99:16899-16903 (2002).
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InterPro; IPR003597; Ig-21.

InterPro; IPR003597; Ig-21.

InterPro; IPR003597; Ig-MIC.

InterPro; IPR003596; Ig-MIC.

InterPro; IPR003596; Ig-V.

INTERPOOR INTERPO
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Best Local
                                                                                   QKIFO MOUSE PRELIMINARY; PRT; 112 AA.

QBKIFO;

QBKIFO;

QBCT-2002 (TrEMBLrel. 22, Created)

QCT-2002 (TrEMBLrel. 22, Last sequence update)

QBCT-2004 (TrEMBLrel. 26, Last annotation update)

Anti-VIPase light chain variable region (Fragment).

Mus musculus (Mouse).
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Mammalia; Butheria; Buarchontoglires; Glires; Rodentia; Sciurognathi;
Muroidea; Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NIH MGC Project;
Submitted (APR-2005) to the EMBL/GenBank/DDBJ databases.
EMBL; BC092251; AMP32251.1; -; mRNA.
SMR; Q569Y8; 23-237.
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurogna
Muridae; Murinae; Mus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           78.4%;
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Pred. No. 1e-34;
8; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 2; Length 237;
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SEQUENCE
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Zhou Y.-X., Taguchi H., Planque S., Karle S., Nishiyama
Submitted (MAY-2002) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EMBL; AF516285; AAM64203.1; -;
PIR; A33933; A33933.
PIR; PC4405; PC4405.
HSSP; P01837; 10RQ.
SMR; Q8K1P0; 3-112.
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InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
SMART; SM00406; IGv; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       nonmutated germ-line genes.";
Proc. Natl. Acad. Sci. U.S.A.
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"Two murine natural polyreactive autoantibodies are encoded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PubMed=2499887;
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                                                                                                                                                                                                                                                                            LTQSPAIMSASPGEKVTTTCRASSSVSSSYLGWYQQXPGSSPRLLIYDTSNLASGVPVRF 63
                                                                                                                                                               SGSGSGTSYSLTISRMEAEDAATYYCQQWSSNPLTFGSGTKLELKR 109
                                                                                                   SGSGSGNSYSLTISSTEGEDAATYYCQQWSSNPPTFGGGTKLEIKR
                                                                                                                                                                                                                                     LTQSPAIMSASPGEKVTMTCSASSSV--SYMHWYQQKSGTSPKRWIYDTSKLASGVPARF
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112 AA; 11901 MW; F6644663201AA239 CRC64;
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November
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   21,
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   2005,
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Pred. No. 1.3e-34;
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   12:04:06
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Search completed: Job time: 162.304

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Result
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Perfect score:
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Maximum Match 100%
Listing first 45 summaries
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    Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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  protein search, using sw model
                                                                                                                                                                                                                                                                                                                                                                         Query
Match
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559
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Gapop 10.0 , Gapext 0.5
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  Issued Patents AA:*

| Cgm2_6/ptodata/1/iaa/5_COMB.pep:*
| Cgm2_6/ptodata/1/iaa/6_COMB.pep:*
| Cgm2_6/ptodata/1/iaa/H_COMB.pep:*
| Cgm2_6/ptodata/1/iaa/PGTUS_COMB.pep:*
| Cgm2_6/ptodata/1/iaa/PGTUS_COMB.pep:*
| Cgm2_6/ptodata/1/iaa/backfiles1.pep:*
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Copyright (c) 1993 - 2005 Compugen Ltd.
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US-09-726-219A-237
US-09-196-522-237
US-09-196-522-255
US-09-196-522-255
US-09-196-522-255
US-09-196-523-497A-26
US-08-6477-728-58
US-08-477-728-58
US-08-477-728-78
US-08-477-728-31
US-08-48-270-31
US-09-196-522-243
US-09-196-522-243
US-09-196-522-256
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Sequence 237, App
Sequence 237, App
Sequence 255, App
Sequence 255, App
Sequence 26, App1
Sequence 26, App1
Sequence 58, App1
Sequence 31, App1
Sequence 24, App1
Sequence 24, App1
Sequence 24, App1
Sequence 256, App
Sequence 276, App1
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CURRENT APPLICATION NUMBER: CURRENT PILING DATE: 2000- PRIOR APPLICATION NUMBER: G PRIOR FILING DATE: 1990-07- PRIOR FILING DATE: 1990-10- PRIOR FILING DATE: 1990-10- PRIOR APPLICATION NUMBER: G PRIOR FILING DATE: 1990-11- PRIOR APPLICATION NUMBER: G PRIOR APPLICATION NUMBER: G PRIOR FILING DATE: 1990-11- PRIOR FILING DATE: 1991-03- PRIOR PILING DATE: 1991-03- PRIOR APPLICATION NUMBER: G PRIOR APPLICATION NUMBER: G PRIOR PILING DATE: 1991-07- PRIOR PILING DATE: 1991-07- PRIOR PILING DATE: 1991-07- PRIOR APPLICATION NUMBER: U PRIOR FILING DATE: 1993-01- PRIOR PILING DATE: 1993-01- PRIOR PILING DATE: 1993-01- PRIOR PILING DATE: 1993-01- PRIOR PILING DATE: 1995-66- NUMBER OF SEQ ID NOS: 272 SOFTWARE: Patentin version BQ ID NO 237 LENGTH: 110 TYPE: PRI DRGANISM: Artificial Seque	5-219A-237 NO. 6806079 NO. 6806079 LINFORMATION CANT: Cambrid CANT: Cambrid CANT: Medica CANT: Pope, CANT: Johnso CANT: Hoogen CANT: Hooling CANT: Chiswe CANT: Bonert CANT: Bonert CANT: Bonert CANTINGENTION	463 461 461 461 461 461 461 461 461 461 461
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US/09/726,219A 1-28 1-28 1-28 1-28 1-28 9015198.6 9022845.3 90224503.6 9024503.6 2 9104744.9 96 9110549.4 15 17/GB91/01134 10 10 10 10 10 10 10 10 10 10	ion US/09726219A  Antibody Technology Lantibody Technology Lesearch Council y, John hony Kevin Hendricus M. Hendricus Rasper Mes Primothy Imothy Imothy Imothy Javid For Producing Methods for Produci	US-09-196-522-25 US-08-469-658-57 US-08-469-658-57 US-09-726-219A-2 US-09-726-219A-2 US-09-196-522-25 US-09-196-522-25 US-09-196-522-25 US-09-618-869-8 US-09-618-869-8 US-09-171-945-9
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	of Specific Binding Pair	Sequence 254, Apple Sequence 57, Appl Sequence 251, Apple Sequence 251, Apple Sequence 250, Apple Sequence 27, Apple Sequence 27, Apple Sequence 27, Appli Sequence 8, Appli Sequence 8, Appli Sequence 9, Appli Sequence 9, Appli Sequence 9, Appli Sequence 9, Appli Sequence 1, Appli Sequence 253, Appli Sequence 257, Appli Seque

FEATURE: .
OTHER INFORMATION: VL of scFv from mouse immunized with 2-phenyl-5-oxazolone

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LENGTH: 110
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Patent No. 691660
                                                                     Matches
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APPLICANT: Cambridge Antibody Technology
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APPLICANT: Bonert, Timothy
TITLE OF INVENTION: Methods for Producing Members of Specific Binding Pairs
FILE REFERENCE: 213839-00004
CURRENT FILING DATE: 1998-11-28
PRIOR APPLICATION NUMBER: GB 9015198.6
PRIOR FILING DATE: 1990-07-10
PRIOR FILING DATE: 1990-07-10
PRIOR PILING DATE: 1990-10-19
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR PILING DATE: 1990-10-19
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR PILING DATE: 1990-10-19
PRIOR APPLICATION NUMBER: GB 9022845.3
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PRIOR APPLICATION NUMBER: GB 9022845.3
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SOFTWARE: PatentIn version 3.1
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PRIOR APPLICATION NUMBER: US 08/484,893
PRIOR FILING DATE: 1995-06-07
                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: GB 9110549.4
PRIOR TILING DATE: 1991-05-15
PRIOR APPLICATION NUMBER: PCT/GB91/01134
PRIOR FILING DATE: 1991-07-10
PRIOR PPLICATION NUMBER: US 07/971,857
PRIOR FILING DATE: 100.07/971,857
                                                                                                                                                                          FEATURE:
OTHER INFORMATION: VL of scFv from mouse immunized with 2-phenyl-5-oxazolone
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: GB 9024503.6
FILING DATE: 1990-11-12
APPLICATION NUMBER: GB 9104744.9
FILING DATE: 1991-03-06
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DIELTOSPAIMSASPGEKVTTTCRASSSVSSSYLGMYQQKPGSSPRLLIYDTSNLASGVP
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Marks, James
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Griffiths, Andrew
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                                                                     Conservative
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86.2%;
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                                                                 Score 484; DB 2; Length 110; Pred. No. 7.3e-39; 7; Mismatches 8; Indels
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US-08-553-497A-20
                                                                                       Query Match
Best Local &
                                                                      Matches
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                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO PCT/EP95/00978
FILING DATE: 16-MAR-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: EP 94104160.0
FILING DATE: 17-MAR-1994
PRIOR APPLICATION NUMBER: EP 94118970.6
FILING DATE: 02-DEC-1994
ATTORNEY/AGENT INFORMATION:
NAME: HAMLET-KING, DIANA
REGISTRATION NUMBER: 33,302
REFERENCE/DOCKET NUMBER: MERCK 1726
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IRM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Ver
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/553,497A
FILING DATE: 17-NOV-1995
                                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
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                                                                                                                                                            MOLECULE TYPE: protein
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                                                                                                                                                                                                                                                                                     TELEPHONE:
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                                 1 DIELTOSPAIMSASPGEKVTTTCRASSSVSSSYLGWYQQKPGSSPRLLIYDTSNLASGVP 60
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                                                                                       Similarity
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 DIELTQSPAIMSASPGEKVTMTCSASSSV--SYMYMYQQKPGSSPRLLIYDTSNLASGVP 194
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                                                                      Conservative
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ROSELL, ELISABET
BLASCO, FRANCESC
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ANTI-EGER SINGLE-CHAIN FVS AND ANTI-EGER
ANTIBODIES
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                                                                   Score 482; DB 1;
Pred. No. 2.8e-38;
2; Mismatches 6
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                                                                                                      DB 1; Length 244
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61 VRFSGSGSGTSYSLTISRMEAEDAATYYCQQWSSNP--LTFGSGTKLELK 108

195

VRFSGSGSGTSYSLTISRMEAEDAATYYCQQWSSYPPMYTFGGGTKLEIK 244

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CURRENT APPLICATION NUMBER: US/09/726,219A
CURRENT FILING DATE: 2000-11-28
CURRENT FILING DATE: 2000-11-28
PRIOR APPLICATION NUMBER: GB 90.5198.6
PRIOR FILING DATE: 1990-77-10
PRIOR PILING DATE: 1990-10-19
PRIOR PILING DATE: 1990-10-19
PRIOR PILING DATE: 1990-10-19
PRIOR PILING DATE: 1990-10-19
PRIOR PILING DATE: 1990-11-12
PRIOR PILING DATE: 1990-11-12
PRIOR FILING DATE: 1990-11-12
PRIOR PILING DATE: 1990-11-12
PRIOR PILING DATE: 1991-03-06
PRIOR PILING DATE: 1991-03-06
PRIOR APPLICATION NUMBER: GB 9104744.9
PRIOR APPLICATION NUMBER: GB 910549.4
PRIOR PILING DATE: 1991-05-15
PRIOR PILING DATE: 1991-07-10
RESULT 5
US-09-196-522-255
; Sequence 255, Application US/09196522
; Patent No. 6916605
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US-09-726-219A-255
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Best Local S
Matches 95
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SOPTWARE: PatentIn version 3.1
SEQ ID NO 255
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APPLICANT:
APPLICANT:
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PRIOR APPLICATION NUMBER: US 08/484,893
PRIOR FILING DATE: 1995-06-07
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APPLICANT: Bonert, Timothy
TITLE OF INVENTION: Methods for Producing Members of Specific Binding
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ORGANISM: Artificial Sequence
PEATURE:
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95; Conserv
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                                                                                                                                         ARFSGSGSGTSYSLTISRMEAEDAATYYCOORSSYPLTFGAGTKLEIKR 109
                                                                                                                                                                                                                                  DIELTOSPAIMSASPGEKVTMTCSASSSVSSNYLNWYQQKSGASPKLWIYSTSNLASGVP
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Griffiths, Andrew
Jackson, Ronald
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Pred. No. 1.4e-38;
5; Mismatches 9
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Patent No.

58440

Application US/08553497A

GENERAL INFOF APPLICANT:

INFORMATION:

APPLICANT: APPLICANT:

KETTLEBOROUGH, C. BENDIG, MARY M. ANSELL, KEITH H. GUSSOW, DETLEF

ADAN, JAUME MITJANS, FRANSESC ROSELL, ELISABET

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APPLICANT: Bonert, Timothy
APPLICANT: Bonert, Timothy
FILE OF INVENTION: Methods for Producing Members of Specific Binding Pairs
FILE OF INVENTION: Methods for Producing Members of Specific Binding Pairs
FILE OF INVENTION: Methods for Producing Members of Specific Binding Pairs
FILE OF INVENTION: US/99/196,522
CURRENT FILING DATE: 1998-11-28
FRIOR APPLICATION NUMBER: GB 9015198.6
FRIOR FILING DATE: 1990-07-10
FRIOR FILING DATE: 1990-10-19
FRIOR APPLICATION NUMBER: GB 9022845.3
FRIOR APPLICATION NUMBER: GB 9022845.3
FRIOR FILING DATE: 1990-10-19
FRIOR FILING DATE: 1990-11-12
FRIOR APPLICATION NUMBER: GB 904540.6
FRIOR FILING DATE: 1991-03-06
FRIOR APPLICATION NUMBER: GB 9104744.9
FRIOR APPLICATION NUMBER: GB 9104744.9
FRIOR APPLICATION NUMBER: GB 910549.4
FRIOR APPLICATION NUMBER: GB 9110549.4
FRIOR FILING DATE: 1991-05-15
FRIOR FILING DATE: 1991-05-15
                                                                                                                                                                                                                                                                                                                                                                                       ; FEATURE: ; OTHER INFORMATION: VL of scFv from mouse immunized with 2-phenyl-5-oxazolone US-09-196-522-255
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RESULT 6
US-08-553-497A-26
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Best Local Similarity
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APPLICANT:
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APPLICANT:
APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: PCT/GB91/01134
PRIOR FILING DATE: 1991-07-10
PRIOR PPLICATION NUMBER: US 07/971,857
PRIOR FILING DATE: 1993-01-08
PRIOR APPLICATION NUMBER: US 08/484,893
PRIOR FILING DATE: 1995-06-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Cambridge Antibody Technology
APPLICANT: Cambridge Antibody Technology Limited
APPLICANT: Medical Research Council
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 110
TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                          61 ARPSGSGSGTSYSLTISRMEAEDAATYYCQQRSSYPLTFGAGTKLBIKR 109
                                                                                                                             61 VRFSGSGSGTSYSLTISRMEAEDAATYYCQQWSSNPLTFGSGTKLELKR 109
                                                                                                                                                                                                                                                                                            95,
                                                                                                                                                                                                                           DIBLTQSPAIMGASPGEKVTTTCRASSSVSŚSYLGWYQQKPGSSPRLLIYDTSNLASGVP 60
                                                                                                                                                                                          DIELTQSPAIMSASPGEKVTMTCSASSSVSSNYLNWYQQKSGASPKLWIYSTSNLASGVP
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Griffiths, Andrew
Jackson, Ronald
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        McCafferty, John
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Johnson, Kevin
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Pred. No. 1.4e-38;
                                                                                                                                                                                                                                                                                            Mismatches
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                                                                                                                                                                                                                                                                                                                                            Length 110;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
                                                                                                                                                                                        GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: 703-243-6410
INFORMATION FOR SEQ ID NO: 26:
SEQUENCE CHARACTERISTICS:
LENGITH: 242 amino acids
                                                                                            APPLICANT:
                                                                                                                                 APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 17-MAR-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: EP 94118970.6
FILING DATE: 02-DEC-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION: 530
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: WO PCT/EP95/00978
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: PIULATS, JAUME
TITLE OF INVENTION: ANTI-EGFR SINGLE-CHAIN FVS AND ANTI-EGFR
TITLE OF INVENTION: ANTIBODIES
                                       APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: WO POSITION DATE: 16-MAR-1995
PRIOR APPLICATION DATA:
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                                                                                                                                                                      APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: HAMLET-KING, DIANA
REGISTRATION NUMBER: 33,302
REFERENCE/DOCKET NUMBER: MERCK 1726
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/01
FILING DATE: 17-NOV-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                        195 VRFSGSGSGTSYSLTISRMEAEDAATYYCQQWSSYPLTFGAGTKLEIK 242
                                                                                                                                                                                                                                                                                                                                                                                                137 DIELTQSPTIMSASPGEKVTMTCSDSSSVSYTY--WYQQKTGSSPRLLIYDTSNLASGVP 194
                                                                                                                                                                                                                                                                                                                                               61 VRFSGSGSGTSYSLTISRMEAEDAATYYCQQWSSNPLTFGSGTKLELK 108
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   INVENTION:
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2200 CLARENDON BLVD. SUITE 1400
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                                                     ADAN, JAUME
MITJANS, FRANSESC
ROSELL, ELISABET
                                                                                                             GUSSOW, DETLEP
                                     ROSELL, ELISABET
BLASCO, FRANCESC
                                                                                                                               ANSELL,
                                                                                                                                                                      KETTLEBOROUGH, C.
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                                                                                                                                 KEITH H.
                                                                                                                                                       MARY M.
JAUME
ANTI-EGFR SINGLE-CHAIN FVS AND ANTI-EGFR
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Pred. No. 3.5e-38;
"" wismatches 7;
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US-07-634-278-58
; Sequence 58, Application US/07634278
; Patent No. 5530101
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                                                                                                                                            GENERAL INFORMATION:
APPLICANT: QUEEN,
APPLICANT: COMMAN
APPLICANT: SCHWEIDI
APPLICANT: LANDOLF:
APPLICANT: COELING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: WO PCT
FILING DATE: 16-MAR-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: BP 941
FILING DATE: 17-MAR-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: BP 941
FILING DATE: 02-DEC-1994
ATTORNEY AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION: 530 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: ANTIBODIES NUMBER OF SEQUENCES: 30 CORRESPONDENCE ADDRESS:
                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend Khourie and Crew
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION: TELEPHONE: 703-243-6333
                                                                                         IITLE OF INVENTION: IMPROVED HUMANIZED IMMUNOGLOBLINS NUMBER OF SEQUENCES: 113
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: HAMLET-KING, DIANA
REGISTRATION NUMBER: 33,
REFERENCE/DOCKET NUMBER:
                    STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/0 FILING DATE: 17-NOV-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                       195 VRFSGSGSGTSYSLTISRMEAEDAATYYCQQWSSYPLTFGAGTKLEIK 242
                                                                                                                                                                                                                                                                                                                                                                                        61 VRFSGSGSGTSYSLTISRMEAEDAATYYCQQMSSNPLTFGSGTKLELK 108
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Palo Alto
California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             amino acid
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                                                                                                                             SELICK, Harold E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                            SCHNEIDER, William P. LANDOLFI, Nicholas F. COBLINGH, Kathleen L.
                                                                                                                                                                                CO, Man Sung
SCHNEIDER, William
                                                                                                                                                                                                                      QUEEN, Cary L.
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GENERAL INFORMATION:
APPLICANT: QUEEN, CATY L.
APPLICANT: SCHNEIDER, William P.
APPLICANT: SELICK, Harold E.
ITILE OF INVENTION: IMPROVED HUMANIZED IMMUNOGLOBLINS
NUMBER OF SEQUENCES: 113.
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, 8th Floor
CITY: Palo Alto
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 9
US-08-477-728-58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-07-634-278-58
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APPLICATION NUMBER: US 07/310,252
PRIOR APPLICATION NUMBER: US 07/310,252
PRILOR DATE: 13-FEB-1989
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/290,975
PILING DATE: 28-DEC-1988
ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M
REGISTRATION NUMBER: 30,223
REFREENCE/DOCKET NUMBER: 31823-002600
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: (415) 326-2422
INFORMATION FOR SEQ ID NO: 58:
SEQUENCE CHARACTERISTICS:
LENGTH: 106 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PATEN: PC-DOS/MS-DOS
SOFTMARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION DATA:
APPLICATION UNMERR: US/07/634,278
FILING DATB: 19-DEC-1990
CLASSIFICATION: 424
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/590,274
                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                 SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: US
ZIP: 94301
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TOPOLOGY: lir
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            85.2%;
Local Similarity 88.8%;
es 95; Conservation
                                                                                                                                                                                     COUNTRY:
  APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            62
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                                                                                                                                                               9411
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             IVLTQSPAIMSASPGEKVIMICSGSSSVSFMY--WYQQRPGSSPRLLIYDISNLASGVPV 59
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US/08/477,728
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US-08-474-040-58
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US-08-477-728-58
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APPLICATION NUMBER: US 07/634,278
FILING DATE: 19-DEC-1990
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/590,274
FILING DATE: 28-SEP-1990
PRIOR APPLICATION NUMBER: US 07/310,252
APPLICATION NUMBER: US 07/310,252
                        ZIP: 94301
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/474,040
FILING DATE: 07-UN-1995
CLASSIFICATION: 536
CLASSIFICATION: 536
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 58,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (415) 326-2422
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
LENGTH: 106 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 13-FEB-1989
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/290
FILING DATE: 28-DEC-1988
ATTORNEY, AGENT INFORMATION:
NAME: Smith, William M
REGISTRATION NUMBER: 30,223
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: QUEEN, Cary L.
APPLICANT: CO, Man Sung
APPLICANT: SCHNEIDER, William P.
APPLICANT: LANDOLFI, Nicholas F.
APPLICANT: COELINGH, Kathleen L.
APPLICANT: SELICK, Harold B.
TITLE OF INVENTION: IMPROVED HUMANIZED IMMUNOGLOBLINS
NUMBER OF SEQUENCES: 113
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: 11823-002600
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 326-2400
                                                                                                                                                                                                                                                                                     STREET: 379 Lytton Avenue CITY: Palo Alto STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
STRANDEDNESS: sir
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local
                                                                                                                                                                                                                                                             COUNTRY:
                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                62 RFSGSGSGTSYSLTISRMEAEDAATYYCQQWSSNPLTFGSGTKLELK 108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              95;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Application US/08474040
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 Mismatches

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Indels

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Gaps

FILING DATE:

US 07/310,252

APPLICATION NUMBER: US

FILING DATE: APPLICATION NUMBER:

19-DEC-1990 28-SEP-1990

US 07/634,278

07/590,274

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                                                                                                                                                                                                                                                                                                                                                                                                                                                     PATENT NO. 5693762

GENERAL INFORMATION:
APPLICANT: QUEEN, C
APPLICANT: CO, Man
APPLICANT: SCHNEIDE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-08-474-040-58
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Best Local S
            APPLICATION NUMBER: US/08/487,200
ELING DATE: 7-JUN-1995
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/634,278
EILING DATE: 19-DEC-1990
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: (415) 326-2422
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/290,975
FILING DATE: 28-DEC-1988
ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M
REGISTRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 11823-002600
                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew
                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: SELICK, Harold E.
TITLE OF INVENTION: IMPROVED HUMANIZED IMMUNOGLOBLINS
NUMBER OF SEQUENCES: 113
                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE: peptide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                              COUNTRY:
                                                                                                                                                                                                                                                                                              CITY: Palo Alto
                                                                                                                                                                                                                                                                                                               STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRANDEDNESS: single TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ENGTH:
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5693762
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                                                                                                                                                                                                                                                                            California
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CO, Man Sung
SCHNEIDER, William P.
LANDOLFI, Nicholas F.
COELINGH, Kathleen L.
                                                                                                                                                                                                                                                                                                               379 Lytton Avenue
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US 07/590,274
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US-08-484-537-58
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 326-2400
TELEFAX: (415) 326-2422
INFORMATION FOR SEQ ID NO: 58:
SEQUENCE CHARACTERISTICS:
LENGTH: 106 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 58, April 500 No. 6180370
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APPLICANT:
APPLICATION NUMBER: US/07/634,278
FILLING DATE: 19-DEC-1990
APPLICATION NUMBER: US 07/590,274
FILLING DATE: 28-SEP-1990
PRIOR APPLICATION NUMBER: US 07/310,252
FILLING DATE: 13-FEB-1989
                                                                                                 CLASSIFICATION:
CLASSIFICATION DATA:
PRIOR APPLICATION NUMBER: US/07/634,278
                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Floppy disk
COMPUTER: LEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/484,537
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY JAGENT INFORMATION:
NAME: Smith, William M
REGISTRATION NUMBER: 30,223
                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 113
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend Khoutie and Crew
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: QUEEN, Cary L. APPLICANT: CO, Man Sung
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 28-SEP-1990
PRIOR APPLICATION DATA:
APPLICATION UMMBER: US O'
FILING DATE: 13-FEB-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: SELICK, Harold TITLE OF INVENTION: IMPROV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE:
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                                                                                                                                                                   FILING DATE:
                                                                                                                                                                                                                                                                                                                                    COUNTRY:
                                                                                                                                                                                                                                                                                                                                                       STATE:
                                                                                                                                                                                                                                                                                                                                                                          CITY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRANDEDNESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             60 RESGSGSGTSYSLTISRMEAEDAATYYCQQWSTYPLTFGAGTKLELK 106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              62 RESGSGSGTSYSLTISRMEAEDAATYYCQQWSSNPLTEGSGTKLELK 108
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                                                                                                                                                                                                                                                                                                                    94301
                                                                                                                                                                                                                                                                                                                                                   Palo Alto
: California
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Application US/08484537
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SCHNEIDER, William P.
LANDOLFI, Nicholas F.
COELINGH, Kathleen L.
                                                                                                                                                                                                                                                                                                                                                                                         379 Lytton Avenue
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  IMPROVED HUMANIZED IMMUNOGLOBLINS
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Pred. No. 46-38;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE: 28-DEC-

28-DEC-1988

US 07/290,975

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RESULT 13
US-07-634-278-31
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COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOPTWARE: PACENTIN Release #1.0, Version #1.25

SOPTWARE: PACENTIN ELEASE

SOPTWARE: PACENTIN ELEASE

SOPTWARE: PACENTIN ELEASE

OPERATION NUMBER: US/07/634,278

PILING DATE: 19-DEC-1990

CLASSIFICATION DATA:

APPLICATION NUMBER: US 07/590,274

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 07/590,274

FILING DATE: 28-SEP-1990

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 07/310,252

PILING DATE: 13-PEB-1989

PRIOR APPLICATION NUMBER: US 07/290,975

FILING DATE: 28-DEC-1988

ATTORNEY/ACPUT THEORMATION.
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INFORMATION FOR SEQ ID NO: 58:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: QUBEN, Cary L.
APPLICANT: CO, Man Sung
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M
REGISTRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 11823-002600
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 326-2420
TELEPHAN: (415) 326-2422
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: SELICK, HAROLD E.
TITLE OF INVENTION: IMPROVED HUMANIZED IMMUNOGLOBLINS
NUMBER OF SEQUENCES: 113
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ..wnsend and T. CITY: Palo Alto STATE: Cal'e-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 106 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        60
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Application US/07634278
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SCHNEIDER, William P. LANDOLFI, Nicholas F. COELINGH, Kathleen L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Townsend and Townsend Khourie and Crew
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                85.2%; Score 476; DB 2
88.8%; Pred. No. 4e-38;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 2; Length 106;
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TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
US-07-634-278-31
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Matches
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Best Local Similarity 88.8%;
Matches 95; Conservative
                                           APPLICATION NUMBER: US 07/310,252
FILING DATE: 13-FEB-1989
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/290,975
FILING DATE: 28-DEC-1988
ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M
REGISTRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 11823-002600
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELBFAX: (415) 326-2422
INFORMATION FOR SEQ ID NO: 31:
SEQUENCE CHARACTERISTICS:
LENGTH: 128 amino acids
                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/634,278
PILING DATE: 19-DEC-1990
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/590,274
PILING DATE: 28-SEP-1990
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: 11
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 326-2400
                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 1:
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: SCHNEIDER, Willia APPLICANT: SELICK, Harold B. TITLE OF INVENTION: IMPROVED
TELEPHONE: (415)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, 8th Ploor CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE: 07-JUN CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/0 FILING DATE: 07-JUN-1995
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                              (415) 326-2400
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; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-477-728-31
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US-08-474-040-31
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Best Local Similarity 88.8%;
Matches 95; Conservative
                                                                                                                                                                   CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/634,278
FILING DATE: 19-DEC-1990
APPLICATION NUMBER: US 07/590,274
FILING DATE: 28-SEP-1990
PRIOR APPLICATION NUMBER: US 07/310,252
APPLICATION NUMBER: US 07/310,252
APPLICATION NUMBER: US 07/310,252
APPLICATION NUMBER: US 07/290,975
FILING DATE: 13-FEB-1989
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/290,975
FILING DATE: 28-DEC-1988
ATTONNEY/AGENT INFORMATION:
NAME: Smith, William M
REGISTRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 11823-002600
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                       TELEFAX: (415) 326-2422 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
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APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                  MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: SELICK, Harold E.
TITLE OF INVENTION: IMPROVED HUMANIZED IMMUNOGLOBLINS
NUMBER OF SEQUENCES: 113
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/08/474,040 FILING DATE: 07-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET: 379 Lytton Avenue CITY: Palo Alto STATE: California
                                              TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE:
                                                                                     LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ZIP: 94301
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                                                H: 128 amino acids
amino acid
Gy: 1
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SCHNEIDER, William P.
LANDOLFI, Nicholas F.
COELINGH, Kathleen L.
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Best Local S
Matches 95
     82
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                                                        24 IVLTQSPAIMSASPGEKVTMTCSGSSSVSFMY--WYQQRPGSSPRLLIYDTSNLASGVPV 81
                                                                           2 IELTQSPAINSASPGEKVTTTCRASSSVSSSYLGWYQQKPGSSPRLLIYDTSNLASGVPV
                                                                                                             95; Conservative
                                                                                                                            Similarity
RFSGSGSGTSYSLTISRMEAEDAATYYCQQWSSNPLTFGSGTKLELK 108
                                                                                                             85.2%; Score 476; DB 1;
88.8%; Pred. No. 5%-38;
tive 3; Mismatches
                                                                                                                                         DB 1; Length 128;
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Search completed: November 21, 2005, 12:07:36
Job time : 41.1245 secs

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No.
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Maximum DB
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Maximum Match 100%
Listing first 45 summaries
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Title:
Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           on:
 Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     seq length: 0 seq length: 2000000000
                                                                         protein search, using sw model
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Match Length
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559
 November 21, 2005, 11:50:51; Search time 134.952 Seconds (without alignments) 337.478 Million cell updates/sec
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1: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*

2: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

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5: /cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.pep:*

6: /cgn2_6/ptodata/1/pubpaa/US11_PUBCOMB.pep:*
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US-10-632-706-88
US-10-632-706-87
US-09-144-886-75
US-09-144-886-75
US-10-632-706-75
US-10-632-706-75
US-10-344-779-2
US-10-632-706-72
US-10-632-706-72
US-10-632-706-73
US-10-803-623-237
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US-10-803-623-255
US-10-803-623-255
US-10-468-370-643
US-10-468-370-643
US-10-895-135-47
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Sequence 88, Appl
Sequence 91, Appl
Sequence 91, Appl
Sequence 78, Appl
Sequence 75, Appl
Sequence 75, Appl
Sequence 2, Appli
Sequence 2, Appli
Sequence 2, Appli
Sequence 27, Appl
Sequence 27, Appl
Sequence 29, Appl
Sequence 21, Appl
Sequence 237, App
Sequence 237, App
Sequence 255, App
Sequence 255, App
Sequence 17, Appl
Sequence 18, Appl
Sequence 18, Appl
Sequence 41, Appl
Sequence 43, Appl
Sequence 47, Appl
Sequence 243, Appl
Sequence 243, Appl
Sequence 243, Appl
Sequence 243, Appl
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RESULT 2  US-10-632-706-87  ; Sequence 87, Application publication No. US2004017; GENERAL INFORMATION:  ; APPLICANT: MARKS, JAMES APPLICANT: MARKSOFFER, APPLICANTION: THE TITLE OF INVENTION: NEU FILE REFERENCE: 407T-895; CURRENT APPLICATION NUMB; CURRENT APPLICATION NUMB;		44444444444444444444444444444444444444
2-706-87 3e 87, A ation No INFORM CANT: MA CANT: MA	1 1706-88 ence 88, Application No. US20 RAL INFORMATION: LICANT: MARKS, J. LICANTICATION: MARKS, J. LICANTICAT	44444444444444444444444444444400000000
6-87 7, Application 7, Application 10, US200401; FORWATION: 1: MARKS, JAMES 1: AMERICON: THE INVENTION: NE INVENTION: NE RENCE: 4077-89 PPLICATION NUM	06-88 06-88 06-88 06 Application Unon No. US20040175. NPORMATION: US20040175. TI: MARKS JAMES D TI: NEURITION: NEURITION: NUMBER: FILING DATE: 2002-0 PLICATION NUMBER: 2002-0 PLICATION NUMBER: 1998-0 PLICATION NUMBER: 1998-0 PLICATION STEEL 1998-0 PSEC ID NOS: 278 109 SEC ID NOS: 278 109 SEC ID NOS: 278 109 CONSERVATION: Singl 106-88 109 CONSERVATION: SINGl 109; CONSERVATION: SINGL 101	844.1 844.1 844.1 83.9 83.9 83.9 83.9 83.9
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	NOCLONAL AN NOCLONAL AND NOCL	61-8 #8-1: 03-612-2 03-612-2 89-193-2 89-417-4 8
ANTIBODIES	ANTIBODIES  DB 4; Le 3.5e-40; eB 0; I SYLGWYOOKPG            SYLGWYOOKPG SYLGWYOOKPG	2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2
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Sequence 78, Application US/09144886
Patent No. US20020155114A1
GENERAL INFORMATION:
APPLICANT: Marks, James D
APPLICANT: Amersdorfer, Peter
TITLE OP INVENTION: Therapeutic Monoclonal Antibodies
TITLE OF INVENTION: Botulinum Neurotoxins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/09/144,886
CURRENT FILING DATE: 1998-08-31
NUMBER OF SEQ ID NOS: 98
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 91
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 91, Application U Patent No. US20020155114A1 GENERAL INFORMATION:
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Best Local &
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PRIOR FILING DATE: 2002-08-01
PRIOR APPLICATION NUMBER: US 09/144,806
PRIOR FILING DATE: 1998-08-31
NUMBER OF SEQ ID NOS: 278
SOFTWARE: Patentin version 3.2
SEQ ID NO 87
                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Marks, James D
APPLICANT: Amersdorfer, Peter
TITLE OF INVENTION: Botulinum Neurotoxins
FILE REFERENCE: 2500.117050
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT
ORGANISM: Artificial Sequence
PEATURE:
OTHER INFORMATION: Description of Artificial Sequence: BONT/A clone
OTHER INFORMATION: 3H4 region VL epitope 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT
ORGANISM: Artificial
FEATURE:
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102; Conserv
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ilarity 93.6%;
Conservative
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Pred. No. 1.1e-36;
1; Mismatches 6
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Pred. No. 1.1e-36;
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                      That
                        Neutralize
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US-10-632-706-75
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Sequence 75, Application US/10632706
Publication No. US20040175385A1
GENERAL INFORMATION:
APPLICANT: MARKS, JAMES D.
APPLICANT: AMERSDORFER, DETER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: Therapeutic Monoclonal TITLE OF INVENTION: Botulinum Neurotoxins FILE REFERENCE: 2500,117USO CURRENT APPLICATION NUMBER: US/09/144,886 CURRENT FILING DATE: 1998-08-31 NUMBER OF SEQ ID NOS: 98 SOFTWARE: Patentin Ver. 2.0 SEQ ID NO 75
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
APPLICANT: Marks, James D
APPLICANT: Amersdorfer, Peter
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CURRENT FILLING DATE: 1998-08-31
NUMBER OF SEQ ID NOS: 98
SOFTWARE: Patentin Ver. 2.0
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                                                                                                                                                                                                                                                                                                                                                                                                       OTHER INFORMATION: Description of Artificial Sequence: BONT/A clone OTHER INFORMATION: C15 region VL epitope 1
                                                                                                                                                                                                                                                                                                                                                                                                                                           ORGANISM: Artificial Sequence FEATURE:
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TYPE: PRT
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89.9%; Pred. No. 1,2e-34;
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Pred. No. 3.7e-35;
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TITLE OF INVENTION: THERAPEUTIC MONOCLONAL ANTIBODIES THAT NEUTRALIZE BOTULINUM TITLE OF INVENTION: NEUROTOXINS

FILE REFERENCE: 407T-895120US

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CURRENT APPLICATION NUMBER: US/10/632,706
CURRENT FILING DATE: 2003-08-01
PRIOR APPLICATION NUMBER: US 60/400,721
PRIOR FILING DATE: 2002-08-01
PRIOR APPLICATION NUMBER: US 09/144,806
PRIOR PILING DATE: 1998-08-31
PRIOR FILING DATE: 1998-08-31
UMBER OF SEQ ID NOS: 278
SOFTWARE: Patentin version 3.2
SEQ ID NO 75
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                                                                                                         US-10-344-779-2
                                                                                                                              RESULT 8
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CURRENT APPLICATION NUMBER: US/10/895,135
CURRENT PILING DATE: 2004-07-21
PRIOR APPLICATION NUMBER: 60/488,447
PRIOR FILING DATE: 2003-07-21
NUMBER OF SEQ ID NOS: 63
SOFTWARE: PATENTIN VETSION 3.2
SEQ ID NO 46
TENEMENT
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                                  Sequence 2, Application US/10344779
Publication No. US20030211106A1
GENERAL INFORMATION:
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Best Local Similarity 90.8%;
Matches 99; Conservative
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Best Local Similarity
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APPLICANT: CHUN, Philip
APPLICANT: TAVARES, Daniel
TITLE OF INVENTION: A CA6 ANTIGEN-SPECIFIC CYTOTOXIC CONJUGATE AND METHODS OF USING
TITLE OF INVENTION: THE SAME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: ImmunoGen, Inc. APPLICANT: PAYNE, Gillian
APPLICANT: SWEET, RAYMOND W. APPLICANT: TORNETTA, MARK A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Artificial FEATURE:
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                      RAYMOND W.
                                                                                                                                                                                                                                                                                                                                                                                               87.8%;
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Pred. No. 1.8e-34;
2; Mismatches 6; Indels
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APPLICANT: TRUNEH, ALEMSEGED
APPLICANT: TRUNEH, ALEMSEGED
APPLICANT: TRUNEH, ALEMSEGED
APPLICANT: TRUNEH, ALEMSEGED
APPLICANT: WATTAM, TREVOR A.
ITITLE OF INVENTION: USEFUL IN TREATMENT OF RANK LIGAND MEDIATED DISORDERS
FILE REFERENCE: GP50033
CURRENT APPLICATION NUMBER: US/10/738,809
CURRENT FILING DATE: 2003-12-17
PRIOR APPLICATION NUMBER: US/10/344,779
PRIOR FILING DATE: 2001-08-21
PRIOR FILING DATE: 2001-08-21
PRIOR PILING DATE: 2000-08-21
PRIOR PILING DATE: 2000-08-21
PRIOR APPLICATION NUMBER: US 60/226,524
PRIOR FILING DATE: 2000-08-21
PRIOR APPLICATION NUMBER: US 60/230,639
PRIOR FILING DATE: 2000-09-07
NUMBER: OF SEQ ID NOS: 18
SOSTWARE: FASESEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; LENGTH: 129
; TYPE: PRT
; ORGANISM: Mus musculus
US-10-344-779-2
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US-10-738-809-2
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                                                                                                                                                                                                                                                 ; SEQ ID NO 2
; LENGTH: 129
; TYPE: PRT
; ORGANISM: Mus musculus
US-10-738-809-2
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Publication No. US20040171117A1
GENERAL INFORMATION:
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                                                                                                                                             Query Match 87.5%; Score 489; DB 4; Length 129; Best Local Similarity 90.7%; Pred. No. 3.8e-34; Matches 98; Conservative 3; Mismatches 5; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 87.5%;
Best Local Similarity 90.7%;
Matches 98; Conservative
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APPLICANT: WATTAM, TREVOR A.

TITLE OF INVENTION: USEFUL IN TREATMENT OF RANK LIGAND MEDIATED DISORDERS FILE REFERENCE: GP5003

FILE REFERENCE: GP5003

CURRENT APPLICATION NUMBER: US/10/344,779

CURRENT FILING DATE: 2003-02-17

PRIOR APPLICATION NUMBER: PCT/US01/26161

PRIOR FILING DATE: 2001-08-21

PRIOR PILING DATE: 2001-08-21

PRIOR APPLICATION NUMBER: US 60/226,524

PRIOR APPLICATION NUMBER: US 60/230,639

PRIOR PILING DATE: 2000-09-07

NUMBER OF SEQ ID NOS: 18

SOFTWARE: FastSEQ for Windows Version 4.0
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  RFSGSGSGTSYSLTISRMEAEDAATYYCQQWSSNPLTFGSGTKLELKR 109
                                                                              IBLTQSPAIMSASPGBKVTTTCRASSSVSSSYLGWYQQKPGSSPRLLIYDTSNLASGVPV 61
                                                 IVLTQSPAIMSASPGEKVTMTCSASSSV--SYMYWYQQKPGSSPRLLIYDTSNLASGVPV 59
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Pred. No. 3.8e-34;
3; Mismatches 5
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Sequence 72, Application US/10632706
publication No. US20040175385A1
GENERAL IMPORMATION:
APPLICANT: MARKS, JAMES D.
APPLICANT: MARKS, JAMES D.
APPLICANT: MARKS, JAMES D.
TITLE OF INVENTION: THERAPEUTIC MONOCLONAL ANTIBODIES THE TITLE OF INVENTION: NEUROTOXINS
FILE REFERENCE: 407T-895120US
CURRENT APPLICATION NUMBER: US/10/632,706
CURRENT FILING DATE: 2003-08-01
PRIOR APPLICATION NUMBER: US 60/400,721
PRIOR APPLICATION NUMBER: US 60/400,721
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US-10-632-706-72
                                                                                                                                                                                                                                                                                                               APPLICANT: Marks, James D
APPLICANT: Amersdorfer, Peter
APPLICANT: Amersdorfer, Peter
TITLE OF INVENTION: Therapeutic Monoclonal Antibodies That Neutralize
TITLE OF INVENTION: Botulinum Neurotoxins
FILE REFERENCE: 2500.117USO
CURRENT APPLICATION NUMBER: US/09/144,886
CURRENT FILING DATE: 1998-08-31
NUMBER OF SEQ ID NOS: 98
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 90
                                                                                 Matches
                                                                                          Query Match
Best Local Similarity
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PRIOR FILING DATE: 1998-08-31
NUMBER OF SEQ ID NOS: 278
SOFTWARE: PatentIn version 3.2
SEQ ID NO 72
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Best Local Similarity
                                                                                                                                                                          TYPE: PRT
ORGANISM: Artificial Sequence
PEATURE:
OTHER INFORMATION: Description of Artificial Sequence: BONT/A clone
OTHER INFORMATION: 3F4 region VL epitope 2
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ORGANISM: Artificial
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                                                                                 97;
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                            1 DIELTQSPAIMSASPGEKVTTTCRASSSVSSSYLGWYQQKPGSSPRLLIYDTSNLASGVP
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DTELTQSPAIMSASPGEKVTMTCSASSSV--SYMYWYQQKPGSSPRLWIYDTSNLASGVP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Application US/09144886
                                                                                 Conservative
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                                                                                                   86.8%;
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                                                                                                   Score 485; DB 3;
Pred. No. 6.9e-34;
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Pred. No. 5.7e-34;
                                                                                 Mismatches
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                                                                                                                     Length 107
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                                                                                 Indels
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58
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CURRENT APPLICATION NUMBER: US/10/803,622
CURRENT FILING DATE: 2004-03-18
FRIOR APPLICATION NUMBER: GB 9015198.6
PRIOR FILING DATE: 1990-07-10
PRIOR FILING DATE: 1990-07-10
PRIOR PPLICATION NUMBER: GB 9022845.3
PRIOR FILING DATE: 1990-10-19
PRIOR FILING DATE: 1990-10-19
PRIOR FILING DATE: 1990-10-19
PRIOR FILING DATE: 1990-10-19
PRIOR APPLICATION NUMBER: GB 9024503.6
PRIOR FILING DATE: 1990-11-12
PRIOR APPLICATION NUMBER: GB 9024744.9
PRIOR FILING DATE: 1990-11-12
PRIOR FILING DATE: 1991-03-06
PRIOR FILING DATE: 1991-03-06
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US-10-803-622-237
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                                                                                                                                                                         Matches
                                                                                                                                                                                                               Query Match
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APPLICANT:
APPLICANT:
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APPLICANT:
APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Winter, Gregory
APPLICANT: Bonert, Timothy
TITLE OF INVENTION: Methods for Producing Members of Specific Binding Pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Cambridge Antibody Technology
APPLICANT: Cambridge Antibody Technology Limited
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                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 110
                                                                                                                                                                                           Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  R FILING DATE: 1991-07-10
APPLICATION NUMBER: US 07/971,857
FILING DATE: 1993-01-08
APPLICATION NUMBER: US 08/484,893
61 ARFSGSGSGTSYSLTISSVEAEDAATYYCQQYSGYPLTFGAGTKLEIKR 109
                            61 VRFSGSGSGTSYSLTISRMEAEDAATYYCQQWSSNPLTFGSGTKLELKR
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                                                                                                                                                                                             Similarity
                                                                                                             DIELTQSPAIMSASPGEKVTTTCRASSSVSSBYLGWYQQKPGSSPRLLIYDTSNLASGVP
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                                                                                DIELTQSPAIMSASPGEKVTMTCRASSSVSSSYLHWYQQKPGASPKVWIYSTSNLASGVP
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Griffiths, Andrew
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Marks, James
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b. US20040157214A1
                                                                                                                                                                       Conservative
                                                                                                                                                                                      86.6%;
86.2%;
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                                                                                                                                                                                        Score 484; DB 4; Length 110; Pred. No. 8.7e-34;
                                                                                                                                                                       Mismatches
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                                                                                                                                                                       Indels
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RESULT 13 US-10-803-653-237

Sequence 237, Application US/10803653
Publication No. US20040157215A1
GENERAL INFORMATION:

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Winter, Gregory
APPLICANT: Winter, Gregory
APPLICANT: Bonert, Timothy
TITLE OF INVENTION: Methods for Producing Men
FILE REFERENCE: 13839-00013
CURRENT APPLICATION NUMBER: US/10/803,653
CURRENT FILING DATE: 2004-03-18
PRIOR APPLICATION NUMBER: GB 9015198.6
PRIOR FILING DATE: 1990-07-10
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR FILING DATE: 1990-10-19
PRIOR PILING DATE: 1990-10-19
PRIOR PILING DATE: 1990-10-19
PRIOR APPLICATION NUMBER: GB 9024503.6
                                                                                                                                                                                   US-09-144-886-96
                                                                                                                                                                                                              RESULT 14
Sequence 96, Application US/09144886
Patent No. US20020155114A1
GENERAL INFORMATION:
APPLICANT: Marks, James D
APPLICANT: Amersdorfer, Peter
APPLICANT: Amersdorfer, Peter
TITLE OF INVENTION: Therapeutic Monoclonal Antibodies That Neutralize
TITLE OF INVENTION: Botulinum Neurotoxins
FILE REFERENCE: 2500.117USO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 272 . SOFTWARE: PatentIn version 3.1 SEQ ID NO 237 LENGTH: 110
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
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Best Local Similarity
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PRIOR APPLICATION NUMBER: US 07/971,857
PRIOR FILING DATE: 1993-01-08
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APPLICANT: Cambridge Antibody Technology Limited
APPLICANT: Medical Research Council
APPLICANT: McCafferty, John
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PRIOR APPLICATION NUMBER: GB 9104744.9
PRIOR FILING DATE: 1991-03-06
PRIOR APPLICATION NUMBER: GB 9110549.4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR PILING DATE: 1991-05-15
PRIOR APPLICATION NUMBER: PCT/GB91/01134
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
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ORGANISM: Artificial Sequence
PEATURE:
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                                                                                                                                                                                                                                                                                  ARFSGSGSGTSYSLTISSVEAEDAATYYCQQYSGYPLTFGAGTKLBIKR 109
                                                                                                                                                                                                                                                                                                          VRFSGSGSGTSYSLTISRMEAEDAATYYCQQWSSNPLTFGSGTKLELKR 109
                                                                                                                                                                                                                                                                                                                                                                              DIBLTOSPAINSASPGEKVTMTCRASSSVSSSYLHWYQQKPGASPKVWIYSTSNLASGVP
                                                                                                                                                                                                                                                                                                                                                                                                      DIELTQSPAIMSASPGEKVTTTCRASSSVSSSYLGWYQQKPGSSPRLLIYDTSNLASGVP
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                                                                                                                                       Application US/09144886
320020155114A1
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Search completed: November 21, 2005, 12:33:34 Job time : 134.952 Becs
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                                                                                                                                                                                                                                                                                                OTHER INFORMATION: single chain antibody US-10-632-706-93
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US-10-632-706-93
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LENGTH: 109
TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: PatentIn version 3.2 SEQ ID NO 93
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CURRENT FILING DATE: 1998-08-31
NUMBER OF SEQ ID NOS: 98
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: THERAPEUTIC MONOCLONAL ANTIBODIES THAT NEUTRALIZE BOTULINUM TITLE OF INVENTION: NEUROTOXINS FILE REFERENCE: 407T-895120US FILE REFERENCE: 407T-895120US FILE REFERENCE: US/10/632,706 CURRENT APPLICATION NUMBER: US/10/632,706 CURRENT FILING DATE: 2003-08-01 PRIOR APPLICATION NUMBER: US 60/400,721 PRIOR PILING DATE: 2002-08-01
                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: US 09/144,806 PRIOR FILING DATE: 1998-08-31 NUMBER OF SEQ ID NOS: 278
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: MARKS, JAMES D.
APPLICANT: AMERSDORFER, E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION: Description of Artificial Sequence: BONT/A clone OTHER INFORMATION: 1H6 region VL epitope 3
                                                                                                                                                                                                                                                                                                                                                LENGTH: 109
TYPE: PRT
ORGANISM: Artificial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FEATURE:
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Pred. No. 1.6e-33;
6; Mismatches
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Pred. No. 1.6e-33;
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    Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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1: /cgn2_6/ptodata/2/pubpaa/USIO_NEW_PUB.pep:*

2: /cgn2_6/ptodata/2/pubpaa/USO7_NEW_PUB.pep:*

3: /cgn2_6/ptodata/2/pubpaa/USO7_NEW_PUB.pep:*

4: /cgn2_6/ptodata/2/pubpaa/USO9_NEW_PUB.pep:*

5: /cgn2_6/ptodata/2/pubpaa/USO9_NEW_PUB.pep:*

6: /cgn2_6/ptodata/2/pubpaa/USO9_NEW_PUB.pep:*

7: /cgn2_6/ptodata/2/pubpaa/USI1_NEW_PUB.pep:*

8: /cgn2_6/ptodata/2/pubpaa/USO9_NEW_PUB.pep:*
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559
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Copyright (c) 1993 - 2005 Compugen Ltd.
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US-11-172-320-8
US-11-172-320-8
US-11-174-186-7
US-11-174-186-7
US-11-174-186-15
US-11-174-186-16-15
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                Sequence 36, Appl
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Sequence 1, Appli
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Sequence 44, Appli
Sequence 27, Appli
Sequence 31, Appli
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	61.8	61.8	61.8	62.3	62.5	62.5	62.5	62.8	63.4	63.4	63.4	63.8	64.0	64.0	64.2	64.7	66.4	66.5	66.5	66.7
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ALIGNMENTS	US-10-512-184-33	US-10-839-799-53	US-10-839-799-43	US-10-839-799-85	US-10-721-763-35	US-10-839-799-65	US-10-839-799-47	US-11-144-248-38	US-10-512-184-68	US-10-512-184-29	US-10-512-184-31	US-11-144-248-14	US-11-144-248-52	US-11-077-978-1	US-11-174-186-5	US-11-144-248-48	<u>.</u>	US-11-174-186-12	US-11-174-186-11	US-11-174-186-14
	Sequence 33, Appl	•	Sequence 43, Appl	•	•	Sequence 65, Appl	•	•	•		) (u	•	2	; <del>-</del>		4.1	: [	; ;	ţ	

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US-10-512-184-36

US-10-512-184-36

; Sequence 36, Application US/10512184

; Publication No. US20050244901A1

; GENERAL INFORMATION:
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RESULT 2
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FILE REFERENCE: 3581.01US01
CURRENT APPLICATION NUMBER: US/10/512,184
CURRENT PILLING DATE: 2004-10-22
NUMBER OF SEQ ID NOS: 72
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                              Query Match
Best Local S
                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION: Description of Artifidial Sequence: scFv SS2 with OTHER INFORMATION: specificity against Sclerotinia sclerotiorum; OTHER INFORMATION: originates from Mus musculus.
                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 248
TYPE: PRT
ORGANISM: Artificial Sequence
PEATURE:
                                                                199
                                                                                                                                                    141 DIVLFQSPVIMSASPGEKVTMTCSASSSV--NYIYMYQMKSGTSPKRMIPDTSKLASGVP
                                                                                            61 VRFSGSGSGTSYSLTISRMEAEDAATYYCQQWSSNPLTFGSGTKLELKR 109
                                                                                                                                                                                      1 DIELTQSPAIMSASPGEKVTTTCRASSSVS9SYLGWYQQKPGSSPRLLIYDTSNLASGVP
                                                                                                                                                                                                                                         86; Conservative
                                                                                                                                                                                                                                                                Similarity
                                                              VRPSGSGGTSFSLTISSMEAEDIATYYCQQWSSPPLTFGAGTKLELKR 247
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78.9%;
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pred. No. 3.5e-19;
7; Mismatches 14
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US-10-512-184-50

Sequence 50, Application US/10512184

Publication No. US20050244901A1

Publication No. US20050244901A1

GENERAL INFORMATION:

APPLICANT: Fraunhofer Gesellschaft zur Friderung der angewandten Forschung

TITLE OF INVENTION: Antibodies, recombinant antibodies, recombinant

TITLE OF INVENTION: antibody fragments and fusions mediated plant disease

TITLE OF INVENTION: resistance against fungi

FILE REFERENCES: 3581.01US01

FILE REFERENCES: 3581.01US01

FULL REFERENCES: 3581.01US01

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GRIANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: fusion protein
OTHER INFORMATION: comprising the leader peptide - chitinase - linker
TOTHER INFORMATION: - scFv SS2 - cmyc/His6.
US-10-512-184-50
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Sequence 8, Application US/11174186
Publication No. US20050244418A1
GENERAL INFORMATION:
APPLICANT: Gillies, Stephen
APPLICANT: Lo, Kin-Ming
APPLICANT: Qian, Xiugi
TITLE OF INVENTION: Recombinant Tumor Specific Antibody
FILE REFERENCE: LEX-019
                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: Recombinant Tumor Specific Antibody And Use Thereof FILE REFERENCE: LEX-019
CURRENT APPLICATION NUMBER: US/11/174,186
CURRENT FILING DATE: 2005-07-01
PRIOR APPLICATION NUMBER: US 60/288,564
PRIOR FILING DATE: 2001-05-03
NUMBER OF SEQ ID NOS: 42
SOFTWARE: Patentin version 3.3
SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 1, Application US/11174186 Publication No. US20050244418A1 GENERAL INFORMATION:
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NUMBER OF SEQ ID NOS: 72
SOFTWARE: PatentIn Ver. 2
SEQ ID NO 50
LENGTH: 615
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Gillies, Stephen
APPLICANT: Lo, Kin-Ming
APPLICANT: Qian, Xiugi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT
ORGANISM: Artificial
FEATURE:
OTHER INFORMATION: KS VK mouse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               540
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                                                                                                                                                                                                                                                 60
                                                                                                                                                                                                                                                                                       62 RFSGSGSGTSYSLTISRMEAEDAATYYCQQWSSNPLTFGSGTKLELK 108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  61 VRFSGSGSGTSYSLTISRMEAEDAATYYCQQWSSNPLTFGSGTKLELKR 109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 DIEUTOSPAIMSASPGEKVTTTCRASSSVSSSYLGWYQQKPGSSPRLLIYDTSNLASGVP
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                                                                                                                                                                                                                                                                                                                                                                                                                Similarity 79.4
85; Conservative
                                                                                                                                                                                                                                                 RFSGSGSGTSYSLIISSMEAEDAATYYCHQRSGYPYTFGGGTKLEIK 106
                                                                                                                                                                                                                                                                                                                            IELTQSPAIMSASPGEKVTTTCRASSSV8SSYLGWYQQKPGSSPRLLIYDTSNLASGVPV 61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               VRFSGSGSGTSFSLTTSSMEAEDIATYYCOOWSSPPLTFGAGTKLELKR 588
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          76.4%;
nilarity 78.9%;
Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                              4; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 414; DB 7; Length 106; Pred. No. 7.4e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 427; DB 1; Length 615, Pred. No. 4.6e-19; 7; Mismatches 14; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                16; Indels
                       And
                       Use Thereof
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PRIOR APPLICATION NUMBER: US 60/288,564
PRIOR FILING DATE: 2001-05-03
NUMBER OF SEQ ID NOS: 42
SOFTWARE: PATENTIN VERSION 3.3
SEQ ID NO 8
LENGTH: 106
TYPE: PRT
ORGANISM: Artificial
RESULT 6
US-11-172-320-4
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US-11-174-186-42
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                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: Recombinant Tumor Specific Antibody And Use Thereof FILE REFERENCE: LEX-019
CURRENT APPLICATION NUMBER: US/11/174,186
CURRENT FILING DATE: 2005-07-01
PRIOR APPLICATION NUMBER: US 60/288,564
PRIOR FILING DATE: 2001-05-03
NUMBER: OF SEQ ID NOS: 42
SOFTWARE: Patentin version 3.3
SEQ ID NO 42
LENGTH: 213
                                                                                                                                                                                                                                                              Best
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APPLICANT: Gillies, Stephen
APPLICANT: Lo, Kin-Ming
APPLICANT: Qian, Xiugi
                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 42, Application US/11174186 Publication No. US20050244418A1
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Best Local (
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CURRENT FILING DATE: 2005-07-01
                                                                                                                                                                                                                                                                                                                                                                   TYPE: PRT
ORGANISM: Artificial
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FEATURE:
                                                                                                                                                                                                                                                                Local
                                                                             59 ARFSGSGSGTSYSLIISSMEAEDAATYYCHORSGYPYTFGGGTKLEIKR 107
                                                                                                           61 VRFSGSGSGTSYSLTISRMEAEDAATYYCQQWSSNPLTFGSGTKLELKR 109
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                                                                                                                                                                                                   1 DIELTQSPAIMSASPGEKVTTTCRASSSVSSSYLGWYQQKPGSSPRLLIYDTSNLASGVP 60
                                                                                                                                                                                                                                                                Similarity
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                                                                                                                                                           IVLTQSPATLSLSPGERVTLTCSASSSV--SYMLMYQQKPGSSPKPWIFDTSNLASGFP
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                                                                                                                                                                                                                                         72.6%; Score 406; DB 7; Length 213; 75.2%; Pred. No. 3.3e-18; tive 7; Mismatches 18; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  5; Mismatches
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Pred. No. 2.1e-18;
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; Sequence 4, Application US/11172320
; Publication No. US20050244413A1
; GENERAL INFORMATION:
; APPLICANT: Adolf, Guenther
; APPLICANT: Baum, Anke
; APPLICANT: Heider, Karl-Heinz
; TITLE OF INVENTION: Compositions and Methods for Treating Cancer using

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CURRENT APPLICATION NUMBER: US/11/172,320
CURRENT APPLICATION NUMBER: US/10/645,215
PRIOR APPLICATION NUMBER: US/10/645,215
PRIOR FILING DATE: 2003-08-21
PRIOR PELICATION NUMBER: EP 02 018 686.2
PRIOR PELICATION NUMBER: EP 02 018 686.2
PRIOR PELICATION NUMBER: US 60/405,956
PRIOR PELICATION NUMBER: US 60/405,956
PRIOR PILING DATE: August 26, 2002
NUMBER: OF SEQ ID NOS: 9
SOPTWARE: PATENTIN VET: 2.1
SEQ ID NO 4
LENGTH: 213
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; ORGANISM: Artificial Sequence
; PEATURE:
; PEATURE:
; OTHER IMPORMATION: Humanised Murine Antibody BIWA 4 Light Chain
US-11-172-320-4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: Gillies, Stephen
APPLICANT: LO, Kin-Ming
APPLICANT: Qian, Xiugi
TITLE OF INVENTION: Recombinant Tumor Specific Antibody And Use Thereof
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS: 42
SOFTWARE: PatentIn version 3.3
SEQ ID NO 9
LENGTH: 106
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                                                                                                                                                                     Matches
                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILE REFERENCE: LEX-019
CURRENT APPLICATION NUMBER: US/11/174,186
CURRENT FILING DATE: 2005-07-01
PRIOR APPLICATION NUMBER: US 60/288,564
PRIOR FILING DATE: 2001-05-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION:
TITLE OF INVENTION:
FILE REFERENCE: 1/138
                                                                                                                                                                                                                                                                                                        ORGANISM: Artificial FEATURE:
                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity 70.6
                                                                                                                                                                                            Match 71.7%;
Local Similarity 75.0%;
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                                                                                                                                                                       81;
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                                       VRFSGSGSGTSYSLTISRMEAEDAATYYCQQWSSNPLTFGSGTKLELK 108
                                                                                  SIVLTQSPATISISPGERVTLTCSASSSV--SYMLWYQQXPGSSPKDWIFDTSNLASGFP
                                                                                                                            DIELTQSPAIMSASPGEKVTTTCRASSSVSSSYLGWYQQKPGSSPRLLIYDTSNLASGVP 60
ARPSGSGSGTSYSLIISSMEABDAATYYCHQRSGYPYTFGGGTKLEIK 106
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      72.5%; Score 405; DB 7; 70.6%; Pred. No. 3.8e-18;
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                                                                                                                                                                7;
                                                                                                                                                                                            Score 401; DB 7; Length 106; Pred. No. 4e-18;
                                                                                                                                                                          Mismatches
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                                                                                       58
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Sequence 27, Application US/10721763

Publication No. US20050249729A1

GENERAL INFORMATION:
APPLICANT: KIRIN BEER KABUSHIKI KAISHA
TITLE OF INVENTION: ANTI TRAIL-R ANTIBODY
FILE REFERENCE: PH-1573-PCT
CURRENT APPLICATION UNMER: US/10/721,763
CURRENT FILING DATE: 2003-11-26
PRIOR APPLICATION NUMBER: JP2001-150213
PRIOR TILING DATE: 2001-05-18
PRIOR APPLICATION NUMBER: JP2001-243040
PRIOR APPLICATION NUMBER: JP2001-243040
PRIOR APPLICATION NUMBER: JP2001-314489
PRIOR PILING DATE: 2001-08-09
PRIOR PILING DATE: 2001-10-11
NUMBER: OF SEQ ID NOS: 45
COPTEMBER: DESCRIPTION OF SEQ ID NOS: 45
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                                                                                                                                                                                                                                                                                            ; TYPE: PRT; ORGANISM: Homo sapiens US-10-721-763-31
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US-10-721-763-27
                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: KIRIN BEER KABUSHIKI KAISHA
TITLE OF INVENTION: ANTI TRAIL-R ANTIBODY
FILE REFERENCE: PH-1573-PCT
CURRENT APPLICATION NUMBER: US/10/721,763
CURRENT FILING DATE: 2003-11-26
PRIOR APPLICATION NUMBER: JP2001-150213
PRIOR PILING DATE: 2001-05-18
PRIOR APPLICATION NUMBER: JP2001-243040
PRIOR APPLICATION NUMBER: JP2001-243040
PRIOR FILING DATE: 2001-08-09
PRIOR FILING DATE: 2001-10-11
NUMBER OF SEQ ID NOS: 45
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Publication No. US20050249729A1
GENERAL INFORMATION:
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LENGTH: 131
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                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                      Query Match 70.8%; Score 395.9; DB 1; Best Local Similarity 69.7%; Pred. No. 9.2e-18;
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                                                                                                                                                                                                                                                                                                                                                                       ENGTH: 128
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               81 DRFSGSGSGTDFTLTTISRLEPEDFAVYYCOOYGSSPLYTFGQGTKLEIKR 130
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  80
                                                 61 VRFSGSGSGTSYSLTISRMEAEDAATYYCOOWSSNPLTFGSGTKLELKR 109
                                                                                               21 BİVLTQSPATLSLSPGERATLSCRASQSV-9SYLAWYQQKPGQAPRLLİYDASNRATGIP
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                                                                                                                                                   1 DIELTQSPAIMSASPGEKVTTTCRASSSVSSSYLGWYQQKPGSSPRLLIYDTSNLASGVP 60
                                                                                                                                                                                               76; Conservative
  13; Mismatches
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RESULT 8 US-10-721-763-27

US-11-172-320-8

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PRIOR APPLICATION NUMBER: US/10/645,215
PRIOR APPLICATION NUMBER: US/10/645,215
PRIOR APPLICATION NUMBER: US/10/645,215
PRIOR APPLICATION NUMBER: BP 02 018 686.2
PRIOR FILING DATE: August 21, 2002
PRIOR APPLICATION NUMBER: US 60/405,956
PRIOR FILING DATE: August 26, 2002
NUMBER OF SEQ ID NOS: 9
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 8
                                                                                                                                                                                            ; FEATURE:
; OTHER INFORMATION: Vk6 light chain US-11-174-186-7
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Publication No. US20050244418A1
GENERAL INFORMATION:
APPLICANT: Gillies, Stephen
APPLICANT: Lo, Kin-Ming
APPLICANT: Qian, Xiugi
                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: Recombinant Tumor Specific Antibody and Use Thereof FILE REFERENCE: LEX-019

CURRENT APPLICATION NUMBER: US/11/174,186

CURRENT FILING DATE: 2005-07-01

PRIOR APPLICATION NUMBER: US 60/288,564

PRIOR FILING DATE: 2001-05-03

NUMBER OF SEQ ID NOS: 42

SOPTWARE: Patentin version 3.3

SEQ ID NO 7
                                                                                               Matches
                                                                                                                   Query Match
Best Local Similarity
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Best Local 8
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APPLICANT: Baum, Anke
APPLICANT: Heider, Karl-Heinz
TITLE OF INVENTION: Compositions and Methods for Treating Cancer using
TITLE OF INVENTION: Cytotoxic CD44 Antibody Immunoconjugates and
TITLE OF INVENTION: Chemotherapeutic Agents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILE REFERENCE: 1/1383
CURRENT APPLICATION NUMBER: US/11/172,320
CURRENT FILING DATE: 2005-06-30
                                                                                                                                                                                                                                                                                             TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                             ORGANISM: Artificial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OTHER INFORMATION: Humanised Antibody BIWA 8 Light Chain
                                                                                                                                                                                                                                                                                                                 LENGTH:
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                    1 DIELTQSPAIMSASPGEKVTTTCRASSSVSSSYLGWYQQKPGSSPRLLIYDTSNLASGVP
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EIVLTQSPATLSLSPGERVTLTCSASSSV--SYMLWYQQKPGQAPKLLIFDTSNLASGIP 58
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                                                                                                                69.9%; Score 391; DB 7; Length 106; 70.4%; Pred. No. 1.4e-17;
                                                                                               11; Mismatches
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US-10-721-763-19
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US-11-174-186-16
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                                                                                                                                                                US-10-721-763-19
                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: ANTI TRAIL-R ANTIBODY
FILE REFERENCE: PH-1573-PCT
CURRENT APPLICATION NUMBER: US/10/721,763
CURRENT FILING DATE: 2003-11-26
PRIOR APPLICATION NUMBER: JP2001-150213
PRIOR APPLICATION NUMBER: JP2001-243040
PRIOR APPLICATION NUMBER: JP2001-243040
PRIOR APPLICATION NUMBER: JP2001-243040
PRIOR APPLICATION NUMBER: JP2001-314489
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Publication No. US20050249729A1 
GENERAL INFORMATION:
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Matches
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Publication No. US20050244418A1
GENERAL INFORMATION:
APPLICANT: Gillies, Stephen
APPLICANT: Lo, Kin-Ming
APPLICANT: Qian, Xiugi
                                                                       Matches
                                                                                           Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                            PRIOR FILING DATE: 2001-10-11
NUMBER OF SEQ ID NOS: 45
SOFTWARE: PatentIn Ver. 2.1
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CURRENT FILING DATE: 2005-07-01
PRIOR APPLICATION NUMBER: US 60/288,564
PRIOR FILING DATE: 2001-05-03
NUMBER OF SEQ ID NOS: 42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: KIRIN BEER KABUSHIKI KAISHA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: Recombinant Tumor Specific Antibody And Use Thereof FILE REFERENCE: LEX-019
                                                                                                                                                                                 TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OTHER INFORMATION: KS VK mouse
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1 DIBLTQSPAIMSASPGEKYTTTCRASSSVSSSYLGWYQQKPGSSPRLLIYDTSNLASGVP 60
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                                                                     Conservative
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                                                                     69.3%; Score 387.5; DB 1; 67.9%; Pred. No. 2.6e-17; vative 15; Mismatches 19;
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21 EIVLTQSPATLSLSPGERATLSCRASQSV-SSFLAWYQQKPGQAPRLLIYDASNRATGIP 79

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RESULT 14
US-11-174-186-15
Sequence 15, Application US/11174186
Publication No. US20050244418A1
GENERAL INFORMATION:
APPLICANT: Gillies, Stephen
APPLICANT: Lo, Kin-Ming
APPLICANT: Qian, Kiugi
APPLICANT: Qian, Kiugi
TITLE OF INVENTION: Recombinant Tumor Specific Antibody And Use Thereof;
FILE REFERENCE: LEX-019
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US-10-721-763-23
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                                                                                                                                                                                                                                        APPLICANT: KIRIN BEER KABUSHIKI KAISHA
TITLE OF INVENTION: ANTI TRAIL-R ANTIBODY
FILE REFERENCE: PH-1573-PCT
CURRENT APPLICATION NUMBER: US/10/721,763
CURRENT FILING DATE: 2003-11-26
PRIOR APPLICATION NUMBER: JP2001-150213
PRIOR FILING DATE: 2001-05-18
PRIOR FILING DATE: 2001-08-09
PRIOR FILING DATE: 2001-08-09
PRIOR APPLICATION NUMBER: JP2001-314489
PRIOR APPLICATION NUMBER: JP2001-314489
PRIOR FILING DATE: 2001-10-11
                                                                                         NUMBER OF SEQ ID NOS: 45
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 23
LENGTH: 129
TYPE: PRT
ORGANISM: Homo sapiens
US-10-721-763-23
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CURRENT FILING DATE: 2005-07-01
PRIOR APPLICATION NUMBER: US 60/288,564
PRIOR FILING DATE: 2001-05-03
NUMBER OF SEQ ID NOS: 42
SOPTWARE: Patentin version 3.3
SEQ ID NO 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 23, Application US/10721763 Publication No. US20050249729A1 GENERAL INFORMATION:
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Query Match 68.2%; Score 381.5; DB 1; Length 129; Best Local Similarity 67.9%; Pred. No. 5.6e-17; Matches 74; Conservative 13; Mismatches 21; Indels 1
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ORGANISM: Artificial
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ENGTH: 106
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1 DIELTQSPAIMSASPGEKVTTTCRASSSVSSSYLGWYQQKPGSSPRLLIYDTSNLASGVP 60

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Db 21 EIVLTQSPATLSLSPGERATLSCRASQSV-9SFLAWYQQKPGQAPRLLIYDASNRATGSP 79

Qy 61 VRFSGSGSTSYSLTISRMEAEDAATYYCQQWSSNPLTFGSGTKLELKR 109
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80 ARFSGSGSGTDFTLTISSLEPEDFAVYYCQQRSDWPLTFGPGTKVDIKR 128

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## SUMMARIES

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Adw4/082 Light cha	Light	Light		Light		Thyroi	Adr59121 Anti-K88/		Mouse E	Murine		3B10x	P5-3 8	Humar	Aay05363 HBV speci	Mur					7	Aar79864 Anti-EGFR	Adr38687 Mouse lig	Description	

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78.7 78.5 78.5	78.7 78.7 78.7	78.9	79.0	79.9 79.2	81.3 80.1	87.9 87.7	89.2 88.1	90.2 90.1
108 108	109	242	110	109 109	108 122	138 109	110 103	138 245
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AAY44993 AAY97236 AAB82710	AAX59306 ADH59678 ADH59707	ABR62132 ABT91209	AAR21306 AAR21305	ABU63531	AAW07440 AAR25958	ADW47092 ADU17908	ADG25837 AAY05362	ADW47094 AAW25783
Aay944993 Aay97236 Aab82710	Adh59678 Adh59707		Aar21306 Aar21305	Adr38686 Abu63531	Aaw07440 Aar25958	Adu17908	Adg2583 / Aay05362	1 M 12
Variable VEGF anta	CPAM4 VK	Single ch		Anti-cucu	ICAM-1 in	Humanised	HBV speci	Light cha Anti-T3 a

## ALIGNMENTS

## Mouse light chain variable region scrv seqid 89. ADR38687 standard; peptide; 109 AA. 02-DEC-2004 ADR38687; (first entry)

antibacterial; antibody; botulinum neurotoxin type A; BONT/A; BONT/A neutralising epitope; anti-botulinum neurotoxin antibody; toxin neutralisation; botulinum neurotoxin poisoning; mouse; light chain variable region; single chain antibody; scFv.

Мив вр

US2004175385-A1.

09-SEP-2004.

01-AUG-2003; 2003US-00632706.

31-AUG-1998; 98US-00144886. 01-AUG-2002; 2002US-0400721P.

(REGC ) UNIV CALIFORNIA.

Marks JD, Amersdorfer P;

WPI; 2004-652009/63.

New isolated antibody that neutralizes botulinum neurotoxin type A, useful for diagnosing botulism or for treating pathologies associated with botulinum neurotoxin poisoning.

Example 1; SEQ ID NO 89; 110pp; English.

ARBSULT 1
ADR3867
ID ADR38
XX ADR38
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XX IIGH The invention describes an isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a specific clone where (I) binds to and neutralises botulinum neurotoxin type A (BoNT/A). An isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a clone chosen from clone 25, C25, C39, IC6, 3D12, B4, IF3, huC25, A1, A22, WR1(V), WR1(T), 3-1, 3-8, 3-10 and ING1, where (I) binds to and neutralizes botulinum neurotoxin type A (BoNT/A). Also described are: a polypeptide (II) comprising BoNT/A neutralising epitope having an epitope that is

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composition metal composition comprising several anti-
condition metal antibodies, where each antibody is specific for a
different epicope of a botulinum neurotoxin and the combination of
antibodies shows greater toxin neutralisation than the single antibodies
comprising (I); and a kit comprising (I). (I) is useful for neutralising
comprising (I); and a kit comprising (I) is useful for neutralising
BONT/A antibody and for neutralising a botulinum neurotoxin which
comprising (I); and a kit comprising (I) in surplus, where each of (I) is
specific for a different epitope of the botulinum neurotoxin and the
combination of antibodies shows greater toxin neutralisation than the
single antibodies in surplus. (I) is useful for diagnosing the botulism
or for treating pathologies associated with botulinum neurotoxin
cor for treating pathologies associated with botulinum neurotoxin
considered to a mouse light chain variable fragment anti-botulinum toxin
core.
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Matches 109
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     Anti-epidermal growth factor receptor (EGFR) single chain antibodies and
                                                           Claim 4; Page 54; 93pp; English
                                                                                                                                                  Anti-EGFR antibodies and single chain from phage-antibody libraries, useful
                                                                                                                                                                                                                                                                                                                         Kettleborough AC,
Mitjans F, Rosell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17-MAR-1994;
02-DEC-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            16-MAR-1995;
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94EP-00118970.
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E, Blasco
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F, Piulats
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06-MAR-1991;
15-MAY-1991;
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                                                                                               Producing members of specific binding pairs - by expression in recombinant host cells with a secreting replicable genetic display
                                                                                                                                                                                                                   Mccafferty J, Pope AR,
Jackson RH, Holliger KP,
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g3p; binding; adsorption; gene VIII; diverse repertoire;
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                                                                         package.
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                                                                                                                                                                                                                                                                                            CAMBRIDGE ANTIBODY TECHNOLOGY MED RES COUNCIL.
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90GB-00024503.
91GB-00004744.
91GB-00010549.
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/label= CDR3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             entry)
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                                                                                                                                                                                                                      Johnson
Marks
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               D-X-G-X-X motif "
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 548; DB 2;
Pred. No. 1.5e-35;
1; Mismatches 2
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Example 22; Fig

24;

209pp; English

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CC The VK sequence is one of 23 (AAR21286-308) expressed from a single chain CC Pv library. The library produces a diverse repertoire of antibody CC cDNA generated from maNA from mice immunised with phox coupled to chicked Serum albumin. The VH and VL kappa sequences were separately amplified by PCR (AAQ23474-84) and ligated into fdCR12 (AAQ23463) for expression on CC the phage surface as fusions with gene III. The resulting library of CC clones was diverse; 23 hapten binding clones were sequenced revealing 8 CC VH genes (A-H; AAR21264-71) in a variety of pairings with 7 VK genes (a-g; AAR21286-92). Most clones were by "crossing" VH-B with the VK clones sequenced. 14 new partners (AAR21293-308) for the VH-B combinations so a further CC clones sequenced. 14 new partners (AAR21293-308) for VH-B were CC dientified. Nearly all the VK genes were "ox-like"; only f, (from the CC original library) and h, p, q, and r (from the hierarchical library) were CC values found. This suggests that phage bearing scFv fragments having weak affinit- ies can be selected with antigen, probably due to the avidity of the multiple antibody heads on the phage. The different combinations CC could also be isolated on a basis of antigen affinity. See also AAR21260-307, 309-311; AAR22450, 565-581
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local S
                                                                                                                                                                                                                                                                                                                  cell-targeting vector; antiviral; cytostatic; immunostimulant; AIDS; gene therapy; immunization; diagnosis; T cell-associated disease; SC acquired immune deficiency syndrome; severe combined immune deficien
                                                                                                                                                                                                                                                                                                                                                             I lymphocyte; antibody; single chain variable antibody; scFv; human;
                                                                                                                                                                                                                                                                                                                                                                                               SNV-env leader/human 7E4-scFv fusion construct.
                                                                                                                                                                                                                                                                                                                                                                                                                                  11-SEP-2003
25-JUN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAB70843
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 110 AA;
                                                                                                                                                                                                        Key
                                                                                                                                                                                                                                       Chimeric.
                                                                                                                                                                                                                                                       Spleen
                                                                                                                                                        Protein
                                                                                                                                                                                                                                                                     Homo sapiens
                                             27-SEP-1999;
                                                                                                           DE19946142-A1
                                                                                                                                                                                      Protein
                                                                                                                                                                                                                                                                                                     cell lymphoma; fusion construct.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          μ
                                                                                                                                                                                                                                                    necrosis virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ARPSGSGGTSYSLTIGTMEAEDVATYYCQQGSSIPYTFGGGTKLEIKR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ARFSGSGSGTSYSLTIGTMEAEDVATYYCQQGSSIPRTFGGGTKLEIKR 109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DSBLTQSPTTMAASPGBKITTTCSASSSISSNYLHWYQQRPGFSPKLLIYRTSNLASGVP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DIELTQSPTTMAASPGEKITITCSASSSISSNYLHWYQQKPGFSPKLLIYRTSNLASGVP
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                                                                                                                                                                                                                                                                                                                                                                                                                                (revised)
(first entry)
                                              99DE-01046142
                                                                                                                                                          1. .45
/label= SNV-env_leader
                                                                                                                                         46. .315
/label= 7E4-scFv
                                                                                                                                                                                                   Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               protein;
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Pred. No. 5.5e-35;
1; Mismatches 3
                                                                                                                                                                                                                                                                                                                     severe combined immune deficiency
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2; Length 110;
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유
                                                                                                                                                                                                                                                      CT This invention describes a novel cell-targeting vector (A) containing a CC DNA sequence (I) encoding a single-chain variable antibody fragment CC (scFv). The products of the invention have antiviral, cytostatic and cc immunostimulant activity and can be used in gene therapy, immunization CC and diagnosis particularly of T cell-associated diseases, specifically CC deficiency (SCID) or T cell lymphoma. (A) target T cells, independently CC deficiency (SCID) or T cell lymphoma. (A) target T cells, independently CC designated 7A5 encodes a 329 amino acid single-chain variable antibody CC designated 7A5 encodes a 329 amino acid single-chain variable antibody CC (squipe osteosarcoma cells susceptible to spleen necrosis virus CC (SNV)) cells, C8166 (human T lymphocyte) cells, and HeLa (human cervical CC (SNV)) cells. After 48 hours, the cells were stained with X-gal to CC carcinoma) cells. After 48 hours, the cells were stained with X-gal to CC showing the high selectivity for human T cells. This sequence represents CC the SNV-env leader/human 7E4-scFv fusion construct used in the invention. The constructused in the invention.
                                                                                                                                                                  Query Match
Best Local &
                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       of e.g. acquired invariable antibody
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cell-targeting vector selective for T lymphocytes, useful in gene therapy of e.g. acquired immune deficiency syndrome, encodes a single-chain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              N-PSDB; AAF61512.
                                                                                                                                                                                                                             Sequence 315
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (BUND ) BUNDESREPUBLIK DEUT PAUL-EHRLICH-INST
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                                                                                                                                                                      Similarity
                                                                          DIELTQSPTTMAASPGEKITITCSASSSISSNYLHWYQQKPGFSPKLLIYRTSNLASGVP 243
ARFSGSGSGTSYSLTIGTMEAEDVATYYCQQGSSIPYTFGGGTKLEIKR
                       ARFSGSGSGTSYSLTIGTMEAEDVATYYCQQGSSIPRTFGGGTKLEIKR 109
                                                                                                 DSELTQSPTTMAASPGEKITTTCSASSSISŚNYLHWYQQRPGFSPKLLIYRTSNLASGVP
                                                                                                                                                    Conservative
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                                                                                                                                                                  96.1%;
                                                                                                                                                  Score 541; DB 4;
Pred. No. 1.6e-34;
1; Mismatches 3;
                                                                                                                                                      ω
••
                                                                                                                                                                                        Length 315;
                                                                                                                                                      Indels
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ARRESULT 5
ARAZIZOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Murine
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAR21299 standard; protein; 110 AA
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                                                    90. .98
/label= CDR3
                                                                                                                                        /label= CDR2
                                                                                                                                                                                                                                                           CDR1
          D-X-G-X-X motif "
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RESULT 6
AAR21295
ID AAR2
XX AAR2
XX AAR2
XC AAR2
XX 21-M
XX
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                                                                                                                                                                                                                                         吊
                                                                                                                                                                                                                                                                                                                                                                                                      The VK sequence is one of 23 (AAR21286-308) expressed from a single chain CC rv library. The library produces a diverse repertoire of antibody CC fragments specific for 2-phenyl-5-oxazolone (phox). It was prepd. using CC DNA generated from mRNA from mice immunised with phox coupled to chicked Serum albumin. The VH and VL kappa sequences were separately amplified by CC PCR (AAQ23474-84) and ligated into fdCA72 (AAQ23463) for expression on CC the phage surface as fusions with gene III. The resulting library of CC clones was diverse; 23 hapten binding clones were sequenced revealing 8 CC VH genes (A-H; AAR21264-71) in a variety of pairings with 7 VK genes (a-g AR21286-92). Most clones were VH-B combinations so a further CC repertoire. The resulting library was screened for hapten binding and 24 CC clones sequenced. 14 new partners (AAR21293-308) for VH-B were CC identified. Nearly all the VK genes were "ox-like; only f, (from the CC values found. This suggests that phage bearing scFv fragments having weak affinit- ies can be selected with antigen, probably due to the avidity of CC values found. This suggests that phage bearing scFv fragments having weak CC uniting antibody heads on the phage. The different combinations CC could also be isolated on a basis of antigen affinity. See also AAR21260-307, 309-311; AAR22450, 565-581
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Best Local Similarity
Matches 105; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    10-JUL-1990;
19-OCT-1990;
12-NOV-1990;
06-MAR-1991;
15-MAY-1991;
                                                                                                                                                                                                                                                                                                                                                                             Sequence 110 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Producing members of specific binding pairs - by expression in recombinant host cells with a secreting replicable genetic display
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mccafferty J,
Jackson RH, 1
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                          21-MAY-1992
                                                         AAR21295;
                                                                                        AAR21295 standard;
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                                                                                                                                                                                             ARFSGSGSGTSYSLTIGTMEAEDVATYYCQQGSSIPRTFGGGTKLEIKR 109
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                                                                                                                                                                                                                                            DIELTOSPTTMAASPGEKITITCSASSSISSNYLHWYQQKPGFSPKLLIYRTSNLASGVP
                                                                                                                                                                          ARFSGSGSGTSYSLTIGTMEAEDVATYYCQQGSSIPFTFGGGTKLEIKR
                                                                                                                                                                                                                                                                                                             Conservative
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                          (first entry)
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90GB-00022845.
90GB-00024503.
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                                                                                        protein; 110
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96.3%;
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Marks JD;
                                                                                                                                                                                                                                                                                                           1;
                                                                                                                                                                                                                                                                                                         Score 540; DB 2; Length 110; Pred. No. 6.6e-35; Indels 1; Mismatches 3; Indels
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S

Matches Query Match Best Local &

Local Similarity

95.4%;

Score 537; DB 2; Length 110; Pred. No. 1.1e-34; 2; Mismatches 3; Indels

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104; Conservative

1 DSELTQSPTTMAASPGEKITTTCSASSSISSNYLHWYQQRPGFSPKLLIYRTSNLASGVP 60

Sequence 110

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The VK sequence is one of 23 (AAR21286-3be) expressed from a single chain CC Fv library. The library produces a diverbe repertoire of antibody CC fragments specific for 2-phenyl-5-oxazobbne (phox). It was prepd. using CC cDNA generated from more from mice immunised with phox coupled to chicked cc serum albumin. The VH and VL kappa sequences were separately amplified by CC pCR (AAQ23474-84) and ligated into fdCATB (AAQ23463) for expression on CC the phage surface as fusions with generality. The resulting library of CC clones was diverse; 23 hapten binding clbnes were sequenced revealing 8 CC yH genes (A-H; AAR21264-71) in a variety of pairings with 7 Vk genes (a-g ; AAR21286-92). Most clones were VH-B combinations so a further CC clones sequenced. 14 new partners (AAR21291-308) for VH-B were CC clones sequenced. 14 new partners (AAR21291-308) for VH-B were CC identified. Nexrly and h, p, q, and r (from the hierarchical library) were Vkox1 type genes. Of the 24 hierarchical clones, one was of type "j". The CC VH-B Vk-d for phox-cABA was 10 nM, one of the highest values found. CC can be selected with antigen, probably due to the avidity of the multiple can be selected with antigen. The different combinations could also be isolated on a basis of antigen affinity. See also AAR21260-307, 309-311; vc.
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Jackson RH, Holliger KP,
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06-MAR-1991;
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19-OCT-1990;
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MED RES COUNCIL.
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90GB-00022845.
90GB-00024503.
91GB-00004744.
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51. .57
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/label= CDR3
/note= " D-X-
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, Marks JD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  D-X-G-X-X motif "
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ARFSGSGSGTSYSLTIGTMEAEDVATYYCQQGSSIPRTFGGGTKLEIKR

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RESULT 7
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                      The VK sequence is one of seven (AAR21286-92) found to be expressed from a single chain FV library from an immunised mouse. The libra- ry produces a diverse repertoire of antibody fragments specific for 2-phenyl-5- oxazolone (phox). It was prepd. using cDNA generated from mRNA from mice immunised with phox coupled to chicked serum albumin. The VH and VL kappa sequences were separately amplified by PCR (see AAQ23474-84) and ligated into fdCAT2 (see AAQ23463) for ex- pression on the phage surface as fusions with gene III. The result- ing library of clones was diverse. Twenty three hapten binding clones were sequenced revealing eight different VH genes (A-H) (see AAR21264-71) in a variety of pairings with the seven different VB genes (a-g). Of the twenty three clones sequenced, two were of type "C", and were "ox-like" genes. (See Berek et al, Nature 316 412-418, 1985). They contain the DXGXX motif in CDR3, the central Gly
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19-OCT-1990;
12-NOV-1990;
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   of which is needed to create a cavity for phOx. Most of the clones were
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90GB-00022845.
90GB-00024503.
91GB-00004744.
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90. .98
/label= CDR3
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Marks JD;
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RESULT 8
ARR2122 8
ARR2122 AX
ARR22 2X
AX
ARR22 AX
ARR22 AX
ARR22 AX
ARR22 AX
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ARR22 AX
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ARR22 ARR22 AX
ARR22 ARR22 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR24 ARR
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Best Local
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19-OCT-1990;
12-NOV-1990;
06-MAR-1991;
15-MAY-1991;
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                                             Example 22; Fig 24; 209pp; English
                                                                                                                 Producing members of specific binding pairs - by expression
recombinant host cells with a secreting replicable genetic or
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MED RES COUNCIL.
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90GB-00022845.
90GB-00024503.
91GB-00004744.
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91GB-00010549.
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/label= CDR2
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/label= CDR1
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95.4%;
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Marks
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Pred. No. 1.6e-34;
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The VK sequence is one of 23 (AAR21286-308) expressed from a single chain

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Best Local Similarity
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                                   Single-stranded antibody against hepatitis B virus core protein, applicable as (gene) therapeutic agents for treatment of hepatitis viral infections.
                                                                                                                                                                                           WPI; 1999-243623/20.
N-PSDB; AAX33931.
                                                                                                                                                                                                                                                                                                      Hayashi N,
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                                                                                                                                                                                                                                                                                                                                                                                                                              (SUMU )
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                                                                                                                                                                                                                                                                                                          Yamamoto
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Pred. No. 2.8e-34;
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                                                                                                                                                                                                                                                                                      (TSUR/)
(ASAK )
(TORI )
                                                                                                   Novel monoclonal antibody against human high-affinity IgE receptor - and DNA fragment encoding the MAb, for the specific identification of human
                                                                                                                                                                                                                                                                                                                                                                                            22-OCT-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                             22-OCT-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Immunoglobulin E; antibody;
complementarity determining
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human IgE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          25-MAR-2003
04-MAR-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAR79162 standard; peptide; 109 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 297 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     as an agent to gene therapy
                                                                           Fc-epsilon RI.
                                                                                                                                                                                  N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAR79162;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   104;
                                                                                                                                                                                                                                                           ) TSURA T.
) ASAHI BREWERIES LTD.
) TORII YAKUHIN KK.
) NIKKA WHISKEY KK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 DSELTQSPTTMAASPGEKITTTCSASSSISSNYLHWYQQRPGFSPKLLIYRTSNLASGVP
                                                                                                                                                                                  AAQ96289
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ARFSGSGSGTSYSLTIGTMEAEDVATYYCOQGSSIPRIFTFGAGTKLEIKR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    receptor-binding, antibody-related peptide light chain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                               93JP-00264792
                                                                                                                                                                                                                                                                                                                                                                                                                                                93JP-00264792
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /label= CDR3L
/note= "all CDR regions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /label= CDR2L
/note= "all C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note= "all CDR regions
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93.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CDR regions
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Pred. No. 1.1e-33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       are claimed"
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Claim

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; NKG2D; NKG2D receptor complex; cancer; infectious disease; tumour; autoimmune disease; head; neck; stomach; oseophagus; colon; liver; ovary; intrahepatic bile duct; pancreas; lung; larynx; breast; uterus; cervix; prostate; kidney; testis; thyroid; bladder; brain; melanoma; myeloma; Fv; sarcoma; leukaemia; lymphoma; virus; bacterium; fungus; protozoan; DAP10; helminth; cytostatic; antimicrobial; immunomodulatory; 1182D10; 6H7E7; 8G7C10; 6ESA7; 1182D1047-7; 8G7C1047-7; 6ESA7x4-7; P4-3; P4-14; P4-15; P5-3; P5-3; P5-9; P5-10; P5-11; P5-23; 3B10xP4-3; 3B10xP4-14; p5-3 tetramerisation domain; 3B10xP5-2; 3B10xP5-23.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 109
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                                                                                                                                                                  Mayer
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DB; AAS97140.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ARPSGSGSGTSYSLTIGTMEAEDVATYYCQQGSSIPLTFGAGTKLELKR 109
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                                                                                                                                                               Riethmueller G,
Hofmeister R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
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Pred. No. 5.7e-34;
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Multifunctional polypeptides comprising binding sites that specifically

Kufer P, Mayer M,

Riethmueller G, Hofmeister R;

Lutterbuese

**7**0

Borschert

Kischel

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to a multifunctional polypeptide comprising a commain with a binding site that specifically recognises an extracellular composition of the NKG2D receptor complex and a second domain which functions can a receptor or ligand. The polypeptide and its associated composition for the treatment of cancer, infections and/or autoimmune confusions. The cancer may be a tumour of the head and neck, stomach, compositions. The cancer may be a tumour of the head and neck, stomach, complying colon, liver, intrahepatic bile ducts, pancreas, lung, colon, liver, intrahepatic bile ducts, pancreas, lung, colon, breast, ovary, uterus, cervix, prostate, kidney, testis, thyroid, complete or brain, or a melanoma, myeloma, sarcoma, leukaemia or lymphoma. The infectious diseases can be caused by viruses, bacteria, fungi, correctious diseases can be caused by viruses, bacteria, fungi, corrections, Grave's disease, ankylosing spondylitis, acute anterior cuveitis, Goodpasture's syndrome, myasthania gravis, insulin-dependent diabetes mellitus, rheumatoid arthritis, pemphigus vulgaris and contoinmune hepatitis. Sequences AU772875 represent the NKG2D contoinmune hepatitis. Sequences AU772875 represent the NKG2D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local (
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                                                                                                                                                                                                                                                                                                  Human; NKG2D; NKG2D receptor complex; cancer; infectious disease; tumour; autoimmune disease; head; neck; stomach; oesophagus; colon; liver; ovary; intrahepatic bile duct; pancreas; lung; llarynx; breast; uterus; cervix; prostate; kidney; testis; thyroid; bladder; brain; melanoma; myeloma; Fv; sarcoma; leukaemia; lymphoma; virus; badterium; fungus; protozoan; DAP10; helminth; cytostatic; antimicrobial; immunomodulatory; 1182D10; 6H7B7; 8G7C10; 6ESA7; 1182D10x4-7; 8G7C10x4-7; 6ESA7X4-7; P4-2; P4-3; P4-14; P4-15; P5-2; P5-3; P5-9; P5-10; P5-11; P5-23; 3B10xP4-3; 3B10xP4-14; p5-3 tetramerisation domain; 3B10xP5-2; 3B10xP5-23.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 256 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  recognize extracellular groups of the NKG2D receptor complex and domains which function as receptors or ligands, useful for treating cancers and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        3B10xP5-23 bispecific single chain Fv.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAU72874 standard; protein;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     receptor and the polypeptides of the invention
                                                                                                   24-MAR-2000; 2000EP-00106467
                                                                                                                                        26-MAR-2001; 2001WO-EP003414
                                                                                                                                                                                                                        WO200171005-A2
                                                             (KUPB/)
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96.2%;
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Pred. No. 1.9e-33;
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    antibacterial; antibody; botulinum neurotoxin type A; BoNT/A; BoNT/A neutralising epitope; anti-botulinum neurotoxin antibody; toxin neutralisation; botulinum neurotoxin poisoning; mouse;
                 Marks JD,
                                                                                                                                                                                  31-AUG-1998; 98US-00144886.
01-AUG-2002; 2002US-0400721P.
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                                                                                                                                                                                                                                                                                                           01-AUG-2003; 2003US-00632706
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US2004175385-A1
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                                                                                                   (REGC ) UNIV CALIFORNIA.
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                 Amersdorfer P;
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Pred. No. 3.7e-33;
1; Mismatches 3
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XX New isolated antibody that neutralizes botulinum neurotoxin type A, PT useful for diagnosing botulism or for treating pathologies associated PT with botulinum neurotoxin poisoning.

XX Example 1; SEQ ID NO 95; 110pp; English.

XX The invention describes an isolated antibody (I) that specifically binds CC to an epitope specifically bound by an antibody expressed by a specific CC clone where (I) binds to and neutralises botulinum neurotoxin type A (BONT/A). An isolated antibody (I) that specifically binds to an epitope composition (II) that positiops (II) and INGI, where (I) binds to and neutralizes botulinum composition (II) and neutralizes botulinum neurotoxin type A (BONT/A). Also described are: a polypeptide (II) comprising BONT/A neutralising epitope that is specifically bound by an antibody expressed by clones as mentioned in (I) comprising (I); and a composition (III) comprising several anti-composition (II) producing (I); and a composition (III) comprising several anti-comprising (I); and a composition neutralisation than the combination of antibodies howe greater toxin neutralisation than the combination of antibodies comprising (I); and a kit comprising (I). (I) is useful for neutralising CC antibodies shows greater toxin neutralisation than the combination of antibodies associated with (I) in surplus. Where each of (I) is specific for a different epitope of the botulinum neurotoxin and the combination of antibodies in surplus. (I) is useful for diagnosing the botulism or treating pathologies associated with botulinum neurotoxin and the combination of antibodies associated with botulinum neurotoxin acid cepton or diagnosis of botulinum neurotoxin acid cepton or diagnosis of botulism. This is the amino acid cepton.

Sequence of a mouse light chain variable fragment anti-botulinum toxin extends the combined of the combined or diagnosis of botulinum coxin coxin coxin coxin coxin coxin coxin coxin coxin coxin coxin coxin coxin coxin coxin coxin coxin coxin coxin coxin coxin coxin coxin coxin coxin coxin
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문 S 밁 ঠ Matches 101; Query Match Local 61 61 Similarity ARFSGSGSGTSYSLTIGTMEAEDVATYYCQQGSSIPRTFGGGTKLEIKR 109 DSELTQSPTTWAASPGEKITTTCSASSSISSNYLHWYQQRPGFSPKLLIYRTSNLASGVP DIELTOSPTTMAASPGEKITITCSASSSIGSNYLHWYQQKPGFSPKLLIYKTSNLASGVP ARFSGSGSGTSYSLTIGAVEAEDVATYYCQQGSSIPYTFGGGTKLEIKR Conservative 93.3%; Score 525; DB 8; Pred. No. 9.8e-34; 3; Mismatches 5 Length 109; Indels 0; Gaps 60 9 0

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                                                                                                                                                                                                                                                                                                                                                                                  The VK sequence is one of 23 (AAR21286-308) expressed from a single chain CC Fv library. The library produces a diverse repertoire of antibody CC fragments specific for 2-phenyl-5-oxazolone (phOx). It was prepd. using CC DNA generated from mRNA from mice immunised with phOx coupled to chicked serum albumin. The VH and VL kappa sequences were separately amplified by CC PCR (AAQ23474-84) and ligated into fdCRT2 (AAQ23463) for expression on CC the phage surface as fusions with gene III. The resulting library of CC clones was diverse; 23 hapten binding clones were sequenced revealing 8 CC VH genes (A-H; AAR21264-71) in a variety of pairings with 7 VK genes (a-g AR21286-92). Most clones were VH-B combinations so a further CC repertoire. The resulting library was screened for hapten binding and 24 CC clones sequenced. 14 new partners (AAR21293-308) for VH-B were CC vkox1 type genes. Of the 24 hierarchical clones, two were of type "o". CC VKox1 type genes. Of the 24 hierarchical clones, two were of type "o". CC viginal library) and h. p. q. and r. (from the hierarchical library) were CC vkox1 type genes. Of the 24 hierarchical clones, two were of type "o". CC viginal library) and h. p. q. and r. (from the hierarchical library) were CC vkox1 type genes. Of the 24 hierarchical clones, two were of type "o". CC viginal library and h. p. q. and r. (from the hierarchical library) were CC cound. This suggests that phage bearing scPv fragments having weak CC affinit- ies can be selected with antigen, probably due to the avidity of CC cound also be isolated on a basis of antigen affinity. See also AAR21260-CC 307, 309-311; AAR22450, 565-581
                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                  Matches 102;
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19-OCT-1990;
12-NOV-1990;
06-MAR-1991;
15-MAY-1991;
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                                                     AAY17963 standard;
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/note= " D-X
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Marks JD;
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Search completed: November 21, Job time: 147.932 Becs

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                                                                                                                 Matches 101;
                                                                                                                                             Query Match
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 SGSGSGTSYSLTIGTMEAEDVATYYCQQGS9IPLTFGAGTKLEIK 239
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96.2%;
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Pred. No. 4.4e-33;
1; Mismatches 3
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64 SGS 	4 LTC     17 LTC	itc)	C;Keywords: heter F;1-13/Domain: si F;14-123/Product: F;14-107/Domain: F;29-104/Domain: F;108-123/Domain:	A;ACCESSION: 010064 A;Molecule type: mRNA A;Residues: 1-121 <ca A;Cross-references: U C;Superfamily: immuno</ca 	e: Alternat	A; ACCESSION: 002207 A; Molecule type: mRNA A; Residues: 1-123 <lev> A; Cross-references: UNI R; Carroll, W.L.; Starne</lev>	R;Levy, S. submitted to the EME A;Reference number:	a chain es: Mus 30-Sep- sion: SC	<b>L</b>		419 418 418 417 416
GSGTSYSL	OSPTTMAAS           SPTTMAAS	94. Similarity 97. 93; Conservative	rotetramer; ignal seque ; Ig kappa V region ( immunoglob	ype: mRNA 1-121 <car> rences: UNIPA</car>	1607-16 ive V ka ir: JL006	mRNA <lev> s: UNIPA Starnes,</lev>	EMBL Data r: S05267	precursor V-J region musculus (house mouse 1991 #sequence revis: 15269; JL0064; 503847		777773355	4444
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EAEDVATYYCOO            EAEDVATYYCOO	TTTCSASSSISSI             TITCSASSSISSI	7%; Score 533; D .2%; Pred. No. 1.2 1; Mismatches	C;Keywords: heterotetramer; immunoglobulin F;1-13/Domain: signal sequence (fragment) #s F;14-123/Product: Ig kappa chain (fragment) F;14-107/Domain: V region (V-kappa-4) <vre>F;29-104/Domain: Immunoglobulin homology <iw (frag<="" (j-kappa-4)="" domain:="" f;108-123="" j="" region="" td=""><td>A;ACCESSION: ULUVO4 A;MOlecule type: mRNA A;Residues: 1-121 <car> A;Cross-references: UNIPARC:UPI0000176799; C;Superfamily: immunoglobulin V region; imm</car></td><td>J. Exp. Med. 168, 1607-1620, 1988 A;Title: Alternative V kappa gene rearrangeménts A;Reference number: JL0061; MUID:89035985; PMID:</td><td>A;McJecule type: mRNA A;McJecule type: mRNA A;Residues: 1-123 <lev> A;Cross-references: UNIPARC:UPI0000115DED; R;Carroll, W.L.; Starnes, C.O.; Levy, R.; I</lev></td><td>Library, February</td><td>30-</td><td>ALIGNMENTS</td><td></td><td></td></iw></vre>	A;ACCESSION: ULUVO4 A;MOlecule type: mRNA A;Residues: 1-121 <car> A;Cross-references: UNIPARC:UPI0000176799; C;Superfamily: immunoglobulin V region; imm</car>	J. Exp. Med. 168, 1607-1620, 1988 A;Title: Alternative V kappa gene rearrangeménts A;Reference number: JL0061; MUID:89035985; PMID:	A;McJecule type: mRNA A;McJecule type: mRNA A;Residues: 1-123 <lev> A;Cross-references: UNIPARC:UPI0000115DED; R;Carroll, W.L.; Starnes, C.O.; Levy, R.; I</lev>	Library, February	30-	ALIGNMENTS		
SGSGSGTSYSLTIGTMEAEDVATYYCQQGSSIPRTFGGGTKLEIKR 	LTQSPTTMAASPGEKITTTCSASSSISSNYLHWYQQRPGFSPKILIYRTSNLASGVPARP 	B 2; Le e-38; 2;	#status pi #status ment) <i< td=""><td>(14099 bulin</td><td>eménts in a murine PMID:3141553</td><td>(14099;</td><td>cy 1989</td><td>- mouse</td><td>INTS</td><td></td><td>. تواسب</td></i<>	(14099 bulin	eménts in a murine PMID:3141553	(14099;	cy 1989	- mouse	INTS		. تواسب
109   122	LIYRTSN        LIYRTSN	123 <i>;</i> le	redicted <sig> predicted <mat></mat></sig>	homology	ne B cell	NID:g52568;		(fragment) t_change 21		Ig Light Ig kappa Ig kappa Ig light Ig light Ig kappa Ig kappa Ig kappa Ig kappa Ig kappa anti-D-d Ig kappa	
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	P 63   P 76	0 ;			oma. An	PIDN:CAA32261.		8		10 10 10 10 10 10 10 10 10 10 10 10 10 1	V r V r pre pre

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DIG

Ig kappa chain V region (hybridoma NC19-B8G2) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Species: Mus musculus (house mouse)
C;Date: 21-Nov-1993 #sequence\_revision 10-Nov-1995 #text\_change 21-Jan-2000
C;Accession: S13699
C;Accession: S13699
R;Pennell, C.A.; Mercolino, T.J.; Grdina, T.A.; Arnold, L.W.; Haughton, G.; Clarke, S.H.
Bur. J. Immunol. 19, 1289-1295, 1989
Bur. J. Immunol. 19, 1289-1295, 1989
A;Title: Blased immunoglobulin variable region gene expression by Ly-1 B cells due to clarke characteristics.

cl

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A;Accession: $13699
A;Status: preliminary
A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-109 <PEN>
A;Cross-references: UNIPARC:UPI000011377B; EMBL:X53352; NID:g55303;
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;16-91/Domain: immunoglobulin homology <IMM>
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S05267
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A;Residues: 1-124 <LEV'>
A;Residues: 1-124 <LEV'>
A;Residues: 1-124 <LEV'>
A;Crose-references: UNIPARC:UPI0000115DEB; EMBL:X14097; NID:g52565; PIDN:CAA32259.1; PID
A;Carroll, W.L.; Starnes, C.O.; Levy, R.; Levy, S.

J. Exp. Med. 168, 1607-1620, 1988
A;Title: Alternative V kappa gene rearrangements in a murine B cell lymphoma. An explana
A;Reference number: JL0061; MUID:89035985; PMID:3141553
A;Accession: JL0063
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RiLevy, S.
submitted to the EMBL Data Library,
a.Reference number: 805267
                                                               Ig kappa chain V region (38C13.V6) - mouse
C:Species: Mus musculus (house mouse)
C:Date: 31-Mar-1990 #sequence_revision 31-Mar-1990 #text_change
C:Accession: PS0069
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Cross-references: UNIPARC:UPI0000176798; EMBL:X14097
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
E;1-13/Domain: signal sequence (fragment) #status predicted <SIG>
E;14-114/Product: Ig kappa chain (fragment) #status predicted <MA:
F;14-114/Domain: V region (V-kappa-3) <VRE>
F;14-111/Domain: immunoglobulin homology <IMM>
A;Title:
                     R;Levy, S.; Campbell, M.J.; Levy, J. Exp. Med. 170, 1-13, 1989
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A; Residues: 1-123 < CAR>
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C;Date: 30-Sep-1991 #sequence revision
C;Accession: S05267; JL0063; S03845
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local S
Matches 103
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                                                                                                                                                                                                                                                                                     SGSGSGTSYSLTIGTMEAEDVATYYCQQGSSIPR--TFGGGTKLEIKR 109
                                                                                                                                                                                                                                                                                                                                                SGSGSGTSYSLTIGTMEAEXVATYYCQQGSSIPRTFGSGTKLEIKR 109
                                                                                                                                                                                                                                                             SGSGSGTSYSLTIGTMEAEDVATYYCOOGSSIPRGVTFGSGTKLEIKR 124
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ilarity 95.3%;
Conservative :
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immunoglobulin light chain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     92.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 522; DB 2; Length 124; Pred. No. 1.9e-37;
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Pred. No. 1.6e-37;
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genes are replaced
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                                                                                                                                                                                    A;Title: Isolation and structural characterization of trimeric A;Reference number: S66536; MUID:95377318; PMID:7649183 A;Accession: S66536
                                                                                                                                                                                                                                                        C;Accession: S66536
R;Tsiotis, G.; Haase, W.; Engel, A.; Michel,
Eur. J. Biochem. 231, 823-830, 1995
                                                                    A;Cross-references: UNIPARC:UPI00001137B0; EMBL:X88903; NID:g895870; PIDN:CAA61365.1; C;Superfamily: immunoglobulin V region; immunoglobulin homology
                                                                                                                    A; Molecule type: mRNA
A; Residues: 1-120 <TSI>
                                                                                                                                                              A; Status: preliminary
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Query Match

/Domain:

immunoglobulin homology

91.5%;

Score

515; <1MM>

밂 2

Length 120,

cyanobacterial

photosysten

PII

20-Jun-2000

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A;Reference number: A92781; MUID:89310348; PMID:2501443
A;Accession: PS0069
A;Status: translation not shown
A;Molecule type: DNA
A;Residues: 1-108 <LEV>
A;Cross-references: UNIPROT:09JL78; UNIPARC:UPI00001767C7
C;Superfamily: immunoglobulin V region; immunoglobulin hom
C;Keywords: heterotetramer; immunoglobulin
C;Keywords: heterotetramer; immunoglobulin
                                                                                                                                                                                                                                                                                                                                                                                         A,Status: preliminary
A,Molecule type: mRNA
A;Molecule type: mRNA
A;Residues: 1-106 < KAV>
A;Cross-references: UNIPARC:UPI0000113791; EMBL:X59098; NID:g52203;
A;Cross-references: UNIPARC:UPI0000113791; immunbglobulin homology
C;Superfamily: immunoglobulin region; immunbglobulin
C;Keywords: heterotetramer; immunoglobulin
E;16-91/Domain: immunoglobulin homology < IMM>
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$29583

$29583

$4 Appa chain V region - mouse

$C;Species: Mus musculus (house mouse)

$C;Date: 06-Jan-1995 #sequence_revisio
RESULT 6
$66536
$1911ght chain V region - mouse (fragment)
Ig 11ght chain V region - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 28-Oct_1996 #sequence_revision 13-Mar-1997 #text_change
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A;Accession: S29583
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R;Kavaler, J
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Best Local Similarity
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97.1%;
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Pred. No. 2.4e-37;
1; Mismatches 3
                                                                                                                                                                                                                                                                                                            1; Mismatches
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Pred. No. 5, 1e-37;
                                                                                                                                                                                                                                                                                                                                                     Length 106
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PID:

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Ig kappa chain - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 18-Feb.194 #sequence_revision 10-Nov-1995 #text_change 21-Jan-2000
C;Accession: $13695
R;Pennell, C.A.; Mercolino, T.J.; Grdina, T.A.; Arnold, L.W.; Haughton, G.;
R;Pennell, C.A.; Mercolino, T.J.; Grdina, T.A.; Arnold, L.W.; Haughton, G.;
R;Pennell, C.A.; Mercolino, T.J.; Grdina, T.A.; Arnold, L.W.; Haughton, G.;
R;Pennell, C.A.; Muscolino, T.J.; Grdina, T.A.; Arnold, L.W.; Haughton, G.;
R;Pennell, C.A.; Muscolino, T.J.; Grdina, T.A.; Arnold, L.W.; Haughton, G.;
R;Pennell, C.A.; Muscolino, T.J.; Grdina, T.A.; Arnold, L.W.; Haughton, G.;
R;Pennell, C.A.; Muscolino, T.J.; Grdina, T.A.; Arnold, L.W.; Haughton, G.;
R;Pennell, C.A.; Muscolino, T.J.; Grdina, T.A.; Arnold, L.W.; Haughton, G.;
R;Pennell, C.A.; Muscolino, T.J.; Grdina, T.A.; Arnold, L.W.; Haughton, G.;
R;Pennell, C.A.; Muscolino, T.J.; Grdina, T.A.; Arnold, L.W.; Haughton, G.;
R;Pennell, C.A.; Muscolino, T.J.; Grdina, T.A.; Arnold, L.W.; Haughton, G.;
R;Pennell, C.A.; Muscolino, T.J.; Grdina, T.A.; Arnold, L.W.; Haughton, G.;
R;Pennell, C.A.; Muscolino, T.J.; Grdina, T.A.; Arnold, L.W.; Haughton, G.;
R;Pennell, C.A.; Muscolino, T.J.; Grdina, T.A.; Arnold, L.W.; Haughton, G.;
R;Pennell, C.A.; Muscolino, T.J.; Grdina, T.A.; Arnold, L.W.; Haughton, G.;
R;Pennell, C.A.; Muscolino, T.J.; Grdina, T.A.; Arnold, L.W.; Haughton, G.;
R;Pennell, C.A.; Muscolino, T.J.; Grdina, T.A.; Arnold, L.W.; Haughton, G.;
R;Pennell, C.A.; Muscolino, T.J.; Grdina, T.A.; Arnold, L.W.; Haughton, G.;
R;Pennell, C.A.; Muscolino, T.J.; Grdina, T.A.; Arnold, L.W.; Haughton, G.;
R;Pennell, C.A.; Muscolino, T.J.; Grdina, T.A.; Arnold, L.W.; Haughton, G.;
R;Pennell, C.A.; Muscolino, T.J.; Grdina, T.A.; Arnold, L.W.; Haughton, G.;
R;Pennell, C.A.; Muscolino, T.J.; Grdina, T.A.; Arnold, L.W.; Haughton, G.;
R;Pennell, C.A.; Muscolino, T.J.; Grdina, T.A.; Arnold, L.W.; Haughton, G.;
R;Pennell, C.A.; Muscolino, T.J.; Grdina, T.A.; Arnold, L.W.; Haughton, G.;
R;Pennell, C.A.; Muscolino, T.J.; Grdina, T.A.; Arnold,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            C;Species: Mus musculus (house mouse)
C;Date: 06-Jan-1995 #sequence_revisio
C;Accession: S29588
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S29588
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A; Accession: S29588
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submitted to the EMBL Data
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                                                                                                                                                                                                                                                                                                                                                                                                                                               A; Residues: 1-102 <KAV>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ig kappa chain V region -
C; Species: Mus musculus ()
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Status: preliminary
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                                                        SGSGSGTSYSLTIGTMEAEDVATYYCQQGSSIPRTFGGG 102
                                                                                                          LTQSPTTMAASPGEKITITCSASSSISSNYLHWYQQKPGFSPKLLIYRTSNLASGVPARF
                                                                                                                                                               LTOSPTTMAASPGEKITTTCSASSSISSNYLHWYQQRPGFSPKLLIYRTSNLASGVPARF
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Conservative
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97.0%;
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T.J.; Grdina, T.A.; Arnold, L.W.; Haughton, G.; Clarke,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 504; DB 2; Li
Pred. No. 5.2e-36;
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2; Mismatches
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Pred. No. 2e-35;
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7;
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                                                                                                                                                                                                                                                                                                                                                                                                                             PIDN: CAA41817.1;
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Ig heavy chain V region - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Species: Mus musculus (house mouse)
C;Date: 18-Feb-1994 #sequence_revision 10-Nov-1995 #text_change 21-Jan-2000
C;Accession: $13696
R;Pennell, C.A.; Mercolino, T.J.; Grdina, T.A.; Arnold, L.W.; Haughton, G.;
Bur. J. Immunol. 19, 1289-1295, 1989
A;Title: Biased immunoglobulin variable region gene expression by Ly-1 B cel
A;Reference number: $13685; MUID:89338557; PMID:2503389
A;Accession: $13696
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C;Species: Mus musculus (house mouse)
C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 21-Jan-2000
C;Accession: S29582
R;Kaveler, J. The man Date Tibrary april 1991
                                                                                                                                                                                                                                                           A;Cross-references: UNIPARC:UPI0000113778; EMBL:X53349; NID:g55297; C;Superfamily: immunoglobulin V region; immunoglobulin homology C;Keywords: heterotetramer; immunoglobulin
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S13696
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Best Local Similarity
Watches 95; Conserve
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A; Residues: 1-102 <KAV>
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                                                                                                                                                                                                                                                                                                                      A; Molecule type: mRNA
A; Residues: 1-101 < PEN>
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Best Local (
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Similarity 95.9%;
93; Conservative
                                                                                                       LTQSPTTMAASPGEKITTTCSASSSISSNYLHWYQQRPGFSPKLLIYRTSNLASGVPARF
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                                                                              LTQSPTTMAASPGEKITITCSASSSISSNYLHWYQQKPGFSXKLLIYRTSNLASGVPARF
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SGSGSGTSYSLTIGTMEAEDVATYYCQQGS9IPLTFG
                         SGSGSGTSYSLTIGTMEAEDVATTYCQQGS91PRTFG 100
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Pred. No. 4.4e-35;
                                                                                                                                                                                 Score 478; DB 2
Pred. No. 8e-34;
                                                                                                                                                            Mismatches
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      100
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Ig heavy chain V region - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 18-Feb-1994 #sequence\_revision 10-Nov-1995 #text\_change 09-Jul-2004
C;Accession: \$13697

RESULT S13697

11

R;Pennell, C.A.; Mercolino, T.J.; Grdina, T.A.; Arnold, L.W.; Haug Eur. J. Immunol. 19, 1289-1295, 1989 A;Title: Biased immunoglobulin variable region gene expression by A;Reference number: S13685; MUID:89338557; PMID:2503389

L.W.; Haughton, G.; Clarke,

Ly-1 B cells

S.H.

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Ig kappa chain V region - mouse
C;Species: Mus musculus (house mouse)
C;Date: 22-Nov-1993 #sequence_revision 10-Nov-1995 #text_change 21-Jan-2000
C;Accession: S17633
R;Clackson, T.; Hoogenboom, H.R.; Griffiths, A.D.; Winter, G.
Mature 352, 624-628, 1991
A;Title: Making antibody fragments using phage display libraries.
A;Reference number: S17633
A;Accession: S17633
A;Status: preliminary
A;Status: preliminary
A;Status: preliminary
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C;Species: Mus musculus (house mouse)
C;Species: Mus musculus (house mouse)
C;Date: 18-Feb-1994 #sequence_revision 10-Nov-1995 #text_change 09-Jul-2004
C;Accession: S13698
C;Accession: S13699
Experimentl, C.A.; Mercolino, T.J.; Grdina, T.A.; Arnold, L.W.; Haughton, G.; Clarke, Bur. J. Immunol. 19, 1289-1295, 1989
Eur. J. Immunol. 19, 1289-1295, 1989
A;Title: Blased immunojlobulin variable region gene expression by Ly-1 B cells due to A;Reference number: S13685; MUID:89338557; PMID:2503389
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A;Molecule type: mRNA
A;Molecule type: mRNA
A;Residues: 1-102 <PEN>
A;Cross-references: UNIPROT: Q9JL78; UNIPARC: UPI00001767B5; EMBL: X53350
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
                                                                                                        A; Molecule type: nucleic acid
A; Residues: 1-93 < CLA>
A; Cross-references: UNIPARC: UPI00001767FA
A; Cross-references: UNIPARC: UPI00001767FA
C; Superfamily: immunoglobulin V region; imm
C; Keywords: heterotetramer; immunoglobulin
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A; Residues: 1-103 < PEN>
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                    Query Match
Best Local Similarity
                                                                                       8-83/Domain:
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Best Local (
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C;Accession: S17631

R;Clackson, T.; Hoogenboom, H.R.; Griffiths, A.D.; Winter, G.

Nature 352, 624-628, 1991

A;Title: Making antibody fragments using phage display libraries.

A;Reference number: S17230; MUID:91326098; PMID:1907718

A;Accession: S17631

A;Accession: S17631
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C;Species: Mus musculus (house mouse)
C;Date: 22-Nov-1993 #sequence_revision 10-Nov-1995 #text_change 09-Jul-2004
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A;Title: Making antibody fragments using phage display libraries.
A;Reference number: S17230; MUID:91326098; PMID:1907718
A;Accession: S17635
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C;Superfamily: immunoglobulin V region; immunoglobulin hor
C;Keywords: heterotetramer; immunoglobulin
F;8-83/Domain: immunoglobulin homology <IMM>
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R InterPro; IPR003596; Ig-v.
R SMART; SM00406; IGv; I.-
PROSITE; P850835; IG LIKE; 1.
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InterPro; IPR00396; Ig-v.
SMART; SM00406; IGv; I-v.
PROSITE; PS50835; IG_LIKE; 1.
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SEQUENCE 114 AA; 12163 MW;
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NUCLEOTIDE SEQUENCE.
STRAIN=BALB/c; TISSUE=Hyperimmunized spleen;
Zhou Y.-X., Taguchi H., Planque S., Karle S.,
Zhou Y.-X. (May-2002) to the EMBL/GenBank/DDBJ
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01-OCT-2002 (TrEMBLrel. 22, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last sequence update)
Anti-VIPase light chain variable region (Fragment).
Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalla; Butheria; Buarchontoglires; Glires; Rodentia; Sciurognathi;
Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE=92381444; PubMed=1512540; DOI=10.1084/jem.176.3.761; Tillman D.M., Jou N.T., Hill R.J., Marion T.N.; "Both IgM and IgG anti-DNA antibodies are the products of clonally selective B cell stimulation in (NZB x NZW)F1 mice."; J. Exp. Med. 176.761-779(1992).
EMBL; AF516284; AAM64202.1; -; Genomic_DNA.
                                                                                                                                                                                                                                                                                                                                                                     HSSP; P01837; 25C8.
SMR; Q8K1F1; 2-114.
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Proc. Natl. Acad. Sci. U.S.A. 86:4624-4628(1989)
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PubMed=2499887;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NCBI_TaxID=10090,
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DSELTQSPTTMAASPGEKITTTCSASSSISSNYLHWYQQRPGFSPKLLIYRTSNLASGVP
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101 AA; 10778 MW;
                                                                           Conservative
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96.0%;
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                                                                                             Score 454; DB 2;
Pred. No. 1.7e-37;
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Pred. No. 1.3e-41;
1; Mismatches 2
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RESULT 4

QRKLF3 MOUSE
QRKLF3 MOUSE PRELIMINARY; PRT; 112 AA.

ID QRKLF3 MOUSE PRELIMINARY; PRT; 112 AA.

AC QRKLF3;

O1-OCT-2002 (TYEMBLrel. 22, Created)

DT 01-OCT-2002 (TYEMBLrel. 26, Last sequence update)

DT 01-MAR-2004 (TYEMBLrel. 26, Last annotation update)

DE Anti-VIPase light chain variable region (Fragment).

Mus musculus (Mouse).

Sukaryota; Metazoa; Choxdata; Craniata; Vertebrata; Euteleostomi;

CC Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;

NCHI_TaxID=10090;

RN [1]

RP NUCLEOTIDE SEQUENCE.
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01-OCT-2002 (TrEMBLrel. 22, Last sequence update)
01-OCT-2002 (TrEMBLrel. 26, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Anti-VIPase light chain variable region (Fragment).
Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;
Muridae; Murinae; Mus.
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SMR; Q8K1F2; 2-112.
Ensembl; ENSMUSG00000064150;
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Proc. Natl. Acad. Sci. U.S.A. 86:4624-46k8(1989).
EMBL, AF516283; AAM64201.1; -; Genomic_DNA.
EMBL, H33932; H33932.
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STRAIN=BALB/c; TISSUE=Hyperimmunized spleen;
STRAIN=BALB/c; TISSUE=Hyperimmunized spleen;
Zhou Y.-X., Taguchi H., Planque S., Karle S., Nishiyama.
Submitted (MAY-2002) to the EMBL/GenBank/DDBJ databases.
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QBK1F2;
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InterPro; IPR003596; Ig_v.
SMART; SM00406; IGv; 1.
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"Two murine natural polyreactive
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112 AA;
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11953 MW; 4716B87FADB543ED CRC64;
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77.1%;
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PROSITE; PS50835; IG L
NON_TER 1 12 112
NON_TER 112 112
SEQUENCE 112 AA; 12
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Tillman D.M., Jou N.T., Hill R.J., Marion T.N.;
"Both IgM and IgG anti-DNA antibodies are the products of clonally selective B cell stimulation in (NZB x NZW)F1 mice.";
J. Exp. Med. 176:761-779(1992).
EMBL; AF516282; AAM64200.1; -; Genomic_DNA.
PIR; PH1085; PH1085.
HSSP; PO1837; 25C8.
SMR; Q8K1F3; 2-112.
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01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Anti-MOG Zl2 variable light chain (Fragment).
Name-Gm1502; Synonyms-anti-MOG kappa;
Mus musculus (Mouse).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Butheria; Euarchontoglires; Glires; Rodent
Muridae; Murinae; Mus.
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PubMed=2427335;
Caton A.J., Brow
"Structural and
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InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
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PubMed=2499887;
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                                                                                                           NUCLEOTIDE SEQUENCE
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                                                                                                                                                                             (OCT-2001)
   Brownlee G.G., and functional
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12017 MW;
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EMBO J. 5:1577-1587(1986).

EMBL; Ay416331; CAC94866.1; -;

PIR; G27887; G27887.

HSSP; P01834; IMIM.

SMR; Q8VDDO; 23-134.

Ensembl; EMSMUSG00000062047; M

MGI; MGI:2686348; Gm1502.
                                                                                                                                                                                                                                                                                                                                              CHAIN
NON TER
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUCLEOTIDE SEQUENCE.
STRAIN-MRI/MpJ-lpr/lpr; TISSUE-Spleen;
MEDLINE=93156722; PubMed=8429833; DOI=10.1016/0161-5890(93)90089-T;
MEDLINE=93156722; PubMed=8429833; DOI=10.1016/0161-5890(93)90089-T;
Takahashi S., Itoh J., Nose M., Ono M., Yamamoto T., Kyogoku M.;
"Cloning and cDNA sequence analysis of dephritogenic monoclonal antibodies derived from an MRL/lpr lupus mouse.";
Mol. Immunol. 30:177-182(1993).
Mol. Immunol. 30:177-182(1993).
EMBL; D14629; BAA03482.1; -; mRNA.
HSSP; D01679; 2FBJ.
HSSP; D01679; 2FBJ.
SMR; Q811C3; 23-131.
ENSEMD1; ENSEMUSG00000058987; Mus musculus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    01-JUN-2003 (TrEMBLrel. 24, 01-JUN-2003 (TrEMBLrel. 24, 01-MAR-2004 (TrEMBLrel. 26, Imunoglobulin gamma-3 kappa Mus musculus (Mouse).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PROSITE, PS50835; IG LIKE; 1.
NON TER 134 134
SEQUENCE 134 AA; 14525 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             InterPro; IPR007i10; Ig-like.
InterPro; IPR003596; Ig-v.
SMART; SM00406; IGv; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eute
Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia;
Muridae; Murinae; Mus.
MCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Q811C3_MOUSE PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       response to a defined antigenic region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PROSITE; PS50835;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 84
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SGSGSGTSYSLTISSMEAEDAATYYCHQRSGYPWTFGGGTKLEIKR
                                                                                                               LTQSPTTWAASPGEKITTTCSASSSISSNYLHWYQQRPGESPKLLIYRTSNLASGVPARF
       SGSGSGTSYSLTIGTMEAEDVATYYCQQGSSIPR-TFGGGTKLBIK 108
                                                                       LTQSPAIMSASPGEKVTMTCRASSSVRSSYLHWYQQKPGSSPKLWIYSTSNLASGVPVRF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                134 134
134 AA; 14525 MW;
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131
131 AA;
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>131
131
14083 MW;
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Last annotation update)
chain precursor (Fragment).
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Pred. No. 8.1e-35;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mus musculus.
                                                                                                                                                                                                                                             Score 417.5; DB 2
Pred. No. 0.8e-34;
                                                                                                                                                                                                                                                                                                                                                                                                                  Potential.
immunoglobullin gamma-3 kappa
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                                                                                                                                                                                                                                                                                                                                                         5E83656994666E9E CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  the influenza virus
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RA Strausberg R.L., Peingold E.A., Grouse L.H., Derge J.G.,
RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Hopkins R.F., Jordan K., Farmer A.A., Rubin G.M., Hong L.,
RA Hopkins R.F., Jordan K., Farmer A.A., Rubin G.M., Hong L.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
Vallalon D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,
RA Villalon D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,
RA Pahey J., Helton B., Ketteman M., Madan A., Rodrigues S., Sanchez A.,
RA Rodriguez A.C., Grimwood J., Schwatz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Schnerch A., Schein J.E., Jones S.J.M., Marra M.A.,
"Generation and initial analysis of more than 15,000 full-length human rand mouse cDNA sequences.",
Proc. Matl Asad. Sci. U.S. A. 99,16899-1699120021.
                                                                                                                                                                                                                           R InterPro; IPR003599; Ig.

R InterPro; IPR003597; Ig-c1.

R InterPro; IPR003597; Ig-c1.

R InterPro; IPR003596; Ig_MHC.

R InterPro; IPR003596; Ig_WHC.

R InterPro; IPR003596; Ig_V:

R InterPro; IPR003596; Ig_V:

R SMART; SM00407; IGC1-set; 1.

R SMART; SM00407; IGC1; 1.

R SMART; SM00406; IGV; 1.

R SMART; SM00406; IGV; 1.

R PROSITE; PS50835; IG_LIKE; 2.

R PROSITE; PS50835; IG_MHC; UNKNOWN 1.

SEQUENCE 237 AA; 25978 MW; A88596AA47FFB932 CRC64;
                                                                                                                                                                     Query Match
Best Local
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Q569Y8 MOUSE PRELIMINARY;
Q569Y8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Submitted (APR-2005) to the EMBL/GenBank/DDBJ databases. EMBL; BC092251; AAH92251.1; -; mRNA. SMR; Q569Y8; 23-237.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;
Muroidea; Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRAIN=FVB/N; TI
NIH MGC Project;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUCLEOTIDE SEQUENCE.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUCLEOTIDE SEQUENCE
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                                                                                                                                                                   Local
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                                                                           26
                                   64
                                                                                                                                                                       Similarity
                                                                       LTQSPTTMAASPGEKITTTCSASSSISSNYLHWYQQRPGFSPKLLIYRTSNLASGVPARF
                       SGSGSGTSYSLTIGTMEAEDVATYYCQQGSSIPRTFGGGTKLEIKR
   SGSGSGTSYSLTISSMEAEDAATYYCQQWNYYPLTFGGGTRLELKR 131
                                                                                                                                                       Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TISSUE=Kidney;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 sci. U.S.A. 99:16899-16903(2002)
                                                                                                                                                                       74,1%;
76.4%;
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Last sequence update)
Last annotation update)
                                                                                                                                                     7;
                                                                                                                                                     Score 417; DB 2
Pred. No. 2e-33;
7; Mismatches
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                                                                                                                                                                                         DB 2; Length 237;
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Best Local S
Matches 83
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ig kappa chain V-IV; region S107B precursor.
Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;
Muroidea; Muridae; Murinae; Mus.
MCBI TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "MAN S.-P., Max E.E., Seidman J.G., Leder P., Scharff M.D.; "Two kappa immunoglobulin genes are expressed in the myeloma S107 "Cell 26:57-66(1981).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This Swiss-Prot entry is copyright. It is produced through a converge the Swiss Institute of Bioinformatics and the EMBL the European Bioinformatics Institute. There are no restrict
                                                                                                                                                                                                                                                                                                                                                                                                                                                              the European Bioinformatics Institute. There are no restrictions ouse as long as its content is in no way modified and this statement
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MISCELLANEOUS: This protein, in which there is a deletion of two amino acids at the V-J recombination site (after position 118), synthesized but not secreted in cells that express and secrete ti
83
                       61
                                                23
                                                                                                 83;
                                                                        ب
                                                                                                             Similarity
                                                               DSELTQSPTTWAASPGEKITTTCSASSSISSWYLHWYQQRPGFSPKLLIYRTSNLASGVP
               ARFSGSGSGTSYSLTIGTMEAEDVATYYCQQGSSIPRTFGGGTKLEIKR 109
                                                ENVLTQSPAIMAASLGQKVTMTCSASSSVSSSYLHWYQQKSGASPKPLIHRTSNLASGVP
ARFSGSGSGTSYSLTISSVEAEDDATYYCOOWSGYP--FGSGTKLEIKR
                                                                                                 Conservative
                                                                                                             73.9%;
76.1%;
                                                                                                                                                                                                                                                                                                   Immunoglobulin V region; Signal.
                                                                                                 10;
                                                                                                            Score 416; DB 1;
Pred. No. 1.2e-33;
                                                                                                                                                                                      Complementarity-determining-3. Framework-4,
                                                                                                                                                                                                             Complementarity-determining-2. Pramework-3.
                                                                                                                                                                                                                                                  Complementarity-determining-1.
                                                                                                                                                   E4BB73072DCF6BE4 CRC64;
                                                                                                 Mismatches
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                                                                                                 14;
                                                                                                                         Length 129;
                                                                                                 Indels
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RESULT 9
Q9U410 MOUSE
ID Q9U410 MOUSE I
AC Q9U410;

PRELIMINARY;

PRT;

106 AA

45

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RESULT 10
QBK1F0 MOUSE
ID QBK1F0 M
AC QBK1F0 M
AC QBK1F0 T
DT 01-QCT-2
DT 01-MAR 2
DR MUSCIECTI
RC STRAIN-E
RL SUDMitte
RN VICLEOTI
RC STRAIN-E
RN VICLEOTI
RP NUCLEOTI
RP NUCLEOTI
RP PUDMEd-2
RA BACCAla
RT TWO mus
RT TOORMUTAL
RT POCE NE
DR EMBL; AI
DR EMBL; AI
DR PIR; AI
DR PIR; AI
DR PIR; PC4
DR PIR; PC4
DR PIR; PC4
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Song X.T., Feng Z.Q., Qiu Z.N., Li Y.Q., Yu X.C., X Huang H.L., Guan X.H.;
"(Cloning and sequence analysis of the light chain gene of monoclonal anti-idiotypic antibody NP30 of
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HSSP; P01679; 2FBJ.
SMR; Q9U410; 4-106.
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Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalla; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurogna
Muridae; Murinae; Mus.
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01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
Monoclonal anti-idiotypic Schiscosoma japonicum antibody
immunoglobulin light chain variable region (Fragment).
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Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Buarchontoglires; Glires; Rodentia; Sciurogna
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-OCT-2002 (TrEMBLrel. 22, Created)
01-OCT-2002 (TrEMBLrel. 22, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Anti-VIPase light chain variable region (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               QBK1F0 MOUSE PRELIMINARY;
                                                                                                Baccala R., Quang T.V., Gilbert N
"Two murine natural polyreactive
nonmutated germ-line genes.";
Proc. Natl. Acad. Sci. U.S.A. 86:
EMBL; AF516285; AAM64203.1; -; Ge
                                                                                                                                                                                                                                                                                                                                                                                        NUCLEOTIDE SEQUENCE.
STRAIN-BALB/c; TISSUE=Hyperimmunized spleen;
STRAIN-BALB/c; TISSUE=Hyperimmunized spleen;
Zhou Y.-X., Taguchi H., Planque S., Karle S.,
Submitted (MAY-2002) to the EMBL/GenBank/DDBJ
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InterPro; IPR003596; Ig_v.
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18:257-259(2000).
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PubMed=2499887;
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PC4405; PC4405.
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                                                                                                                                                                                                                                                         Gilbert M.,
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Pred. No. 3.1e-33;
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RA Strausberg R.L., Feingold E.A., Grouse I.H., Derge J.G.,
RA Klausner R.D., Collins F.S., Wagner L., Shemmen C.M., Schuler G.D.,
RA Altechul S.F. Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Altechul S.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Stapleton M., Soarse M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Brownstein M.J., Usdin T.B., Toshiyuki 9., Carninci P., Prange C.,
RA Brownstein M.J., Usdin T.B., Toshiyuki 9., Carninci P., Willahy S.J.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Raha S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Rahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Rahey J., Helton E., Ketteman M., Badan A., Rodrigues S., Sanchez A.,
RA Rodriguez A.C., Grimwood J., W., Schwitz J., Dickson M.C.,
RA Rodriguez A.C., Grimwood J., Schwitz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schwitz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schwitz J., Myers R.M.,
RA Rahesley R.W., Touchman J.W., Schwitz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schwitz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schwitz J., Myers R.M.,
RA Rahesley R.W., Touchman J.W., Schwitz J., Myers R.M.,
RA Generation and initial analysis of more than 15,000 full-length human RT and mouse cDNA sequences ",
RA Packer R.M., Myers R.M., Myers R.M., Myers R.M., Myers R.M., Myers R.M., Myers R.M., Myers R.M., Myers R.M., Myers R.M., Myers R.M., Myers R.M., Myers R.M., Myers R.M., Myers R.M., Myers R.M., Myers R.M., Myers R.M., Myers R.M., Myers R.M., Myers R.M., Myers R.M., Myers R.M., Myers R.M., Myers R.M., My
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurogna
Muroldea; Muridae; Murinae; Mus.
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InterPro; IPR003596; Ig_v.
SMART; SM00406; IGv; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUCLEOTIDE SEQUENCE
STRAIN-FVB/N; TISSUI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Igk-C protein.
Name=Igk-C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Q58EV6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Q58EV6_MOUSE PRELIMINARY;
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Ensembl; ENSMUSG00000062047; Mus musculus
                                                                                                                         Submitted (MAR-2005) to the EMBL/GenBank/DDBJ EMBL; BC091738; AAH91738.1; -; mRNA. SMR; Q58EV6; 23-235. GO; GO:0003823; F:antigen binding; IEA.
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                                                                                                                                                                                                                                                        NIH MGC Project;
                                                                                                                                                                                                                                                                                            STRAIN=FVB/N;
                                                                                                                                                                                                                                                                                                                       NUCLEOTIDE SEQUENCE
                                                              InterPro;
                                                                                                  InterPro;
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                                                                    IPR007110;
                                                                                                  IPR003599;
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Pred. No. 5.2e-33;
3; Mismatches 18
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KV6F_MOUSE
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Pfam; PF07654; C1-8et; 1.

SMART; SM00409; IG; 2.

SMART; SM004007; IGc1; 1.

SMART; SM00406; IGv; 1.

PROSITE; PS50835; IG LIKE; 2.

PROSITE; PS50835; IG MHC; UNKNOWN 1.

PROSITE; PS00290; IG MHC; UNKNOWN 1.

PROSITE; PS00290; IG MHC; UNKNOWN 1.
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MEDLINE=83771467; PubMed=6877353;

MEDLINE=83271467; PubMed=6877353;

Markham A.F., Milstein C.

Markham A.F., Milstein C.

"mRNA sequences define an unusually restricted IgG resp

phenyloxazolone and its early diversification.";

Nature 304:320-324(1983).

-i- FUNCTION: Anti-2-phenyl oxazolone (PHOX) Antibody.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertel
Mammalia; Eutheria; Euarchontoglires; Glires;
Muroldea; Muridae; Murinae; Mus.
                                                                                                                                                                                                                        Ensembl; ENSMUSG00000062047; Mus
InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
                                                                                                                                                                                                                                                                                EMBL; K00735; AAA38680.1; -; HSSP; P01679; 2FBJ.
                                                                                                                                                                                                                                                                                                                                               This Swiss-Prot entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on it
                                                                                                                                                                                                   PROSITE; PS50835; IG_LIKE; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mus musculus (Mouse)
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13-AUG-1987 (Rel.
10-MAY-2005 (Rel.
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7654; C1-set; 1.
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VV-V region NQ2-17.4.1.
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Score 401; DB 1;
Pred. No. 3.1e-32;
9; Mismatches 16
                                                                                                                                                                                                                                                                                              mRNA.
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Pred. No. 3
                                                                                           Framework-3.
Complementarity-determining-3
Framework-4.
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Complementarity-determining-2.
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lires; Rodentia; Sciurognathi;
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HSSP; Q91W12; 1AY1.
SMR; P44945; 1-108.
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MEDLINE=83271467; PubMed=6877353;
Kaartinen M., Griffiths G.M., Markham A.F., Milstein
"mRNA sequences define an unusually restricted IgG re
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi; Muroidea; Murinae; Mus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NCBI_TaxID=10090;
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Mus musculus (Mouse)
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InterPro; IPR003596; Ig_v.
                                                                                                                                                                                                                                                                                   Ensembl; ENSMUSG00000062686; Mus musculus
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                                  LTQSPTTMAASPGEKITTTCSASSSISSNYLHWYQQRPGFSPKLLIYRTSNLASGVPARF
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7 (Rel. 05, Last sequence :
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15 hain V-VI region NQ2-6.1.
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73.8%;
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                                                                                                                                                                                                                               domain; Immunoglobulin V
                                                                               ; Score 398; DB 1;
; Pred. No. 6.3e-32;
10; Mismatches 14
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By similari
                                                                                                                                                                   Complementarity-determining-3
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RESULT 14
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P04943; 13-AUG-1987 13-AUG-1987 10-MAY-2005

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13-AUG-1987 (Rel. 05, Last sequence update)
10-MAY-2005 (Rel. 47, Last annotation update)
10 kappa chain V-VI region NQ6-8.3:1.
If kappa chain (Nouse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalla; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;
Muroidea; Muridae; Murinae; Mus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUCLEOTIDE SEQUENCE.
MEDLING=83271467; PubMed=6877353;
MEDLING=83271467; PubMed=6877353;
MARXHAM A.F., Milstein
MRNA sequences define an unusually restricted IgG re
phenyloxazolone and its early diversification.";
Nature 304:320-324(1983).
                                                                    P04941;
13-AUG-1987 (Rel. 05, Created)
13-AUG-1987 (Rel. 05, Last sequence update)
10-MAY-2005 (Rel. 47, Last annotation update)
1g kappa chain V-VI region NQ2-48.2.2.
Mus musculus (Mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EMBL; K00740; AAA38685.1; -; mRNA.
HSSP; P01679; 2FBJ.
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                                Bukaryotaj Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;
Muroidea; Muridae; Murinae; Mus.
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InterPro; IPR003596; Ig_v.
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NUCLEOTIDE SEQUENCE.
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pred. No. 9.8e-32;
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SMR; P04941; 1-103.
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Kaartinen M., Griffiths G.M., Markham A.F., Milstein C.;
"mRNA sequences define an unusually restricted IgG responsive and its early diversification.";
Nature 304:320-324(1983)...
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                                                                                                                                                                                                                                                                                                                                                               use as
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                                         LTQSPAINSASPGQKVTMTCSASSSVS--YMHWYQQKSGTSPKRWIYDTSKLASGVPARF
                                                      LTQSPTTWAASPGEKITTTCSASSSISSNYLHWYQQRPGFSPKLLIYRTSNLASGVPARF
                     SGSGSGTSYSLTIGTMEAEDVATYYCOOGSSIPRTFGGGTKLEIKR
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IPR003596; Ig_v.
  SGSGSATSYSLTITSMQAEDAATYYCQQWSSNPLTFGAGTKLXLKR
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Complementarity-determining-3.
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score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.
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219.131 Million cell updates/sec
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1: /cgn2_6/ptodata/1/laa/5_COMB.pep:*

2: /cgn2_6/ptodata/1/laa/6_COMB.pep:*

3: /cgn2_6/ptodata/1/laa/H_COMB.pep:*

4: /cgn2_6/ptodata/1/laa/RE_COMB.pep:*

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6: /cgn2_6/ptodata/1/laa/RE_COMB.pep:*
                                 GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
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              US-08-553-497A-14
US-09-726-219A-247
US-09-726-219A-249
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US-09-726-219A-255
US-09-726-219A-258
US-09-726-219A-238
US-09-726-219A-238
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                            Sequence 14, Appl
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Sequence 245, App
Sequence 258, App
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Sequence 238, App
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Sequence 255, App
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Sequence 2, Appli Sequence 2, Appli Sequence 92, Appl Sequence 242, App Sequence 242, Appli Sequence 2, Appli Sequence 4, Appli Sequence 1, Appli Sequence 17, Appl Sequence 17, Appl	1 US-08-116-7788-2 1 US-08-438-542-2 1 US-08-483-5288-92 1 US-09-126-219A-242 2 US-09-126-522-242 2 US-09-526-738A-4 2 US-09-526-738A-4 2 US-09-798-689-8 2 US-09-798-689-8 2 US-09-798-689-21 3 US-09-798-689-21 4 LIGNMENTS	78.2 129 78.2 129 78.2 129 78.0 108 78.0 258 77.6 235 77.6 235 77.6 235		00000044444 0000000000000
Sequence 257, Sequence 237, Sequence 240, Sequence 240, Sequence 240, Sequence 254, Sequence 254,			450 450 447 4447	33 33 33 4

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APPLICANT: KETTLEBOROUGH, C. A.
APPLICANT: BENDIG, MARY M.
APPLICANT: BENDIG, MARY M.
APPLICANT: BENDIG, MARY M.
APPLICANT: GUSSOW, DETLEF
APPLICANT: GUSSOW, DETLEF
APPLICANT: GUSSOW, DETLEF
APPLICANT: HIJONS, FRANSESC
APPLICANT: HIJONS, PRANSESC
APPLICANT: PIULATS, JAUME
TITLE OF INVENTION: ANTIBODIES
NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSER: MILLEN, WHITE, ZELANO & BRANIGAN, P.C.
STREET: 2200 CLARENDON BLVD. SUITE 1400
CITY: ARLINGTON
STATE: VA
COUNTRY: US
TIP: 22201
COMPUTER REDABLE FORM:
MEDIUM TYPE: Ploppy disk
COMPUTER: LEM PC compatible
OPERATION STATE: VA
COUNTRY: US
SOFTWARE: PATENTIN SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN BATA:
APPLICATION NUMBER: US/08/53,497A
FILING DATE: 17-NOV-1995
CLASSIFICATION NUMBER: WO PCT/SP95/00978
FILING DATE: 17-MAR-1995
PRIOR APPLICATION NUMBER: BP 94104160.0
FILING DATE: 16-MAR-1995
PRIOR APPLICATION NUMBER: BP 94104160.0
FILING DATE: 17-MAR-1994
PRIOR APPLICATION NUMBER: BP 94104160.0
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PRIOR APPLICATION NUMBER: BP 94104160.0
FILING DATE: 17-MAR-1994
PRIOR APPLICATION NUMBER: BP 94104160.0
FILING DATE: 10-MAR-1994
PRIOR APPLICATION NUMBER: BP 94104160.0
FILING DATE: 02-DEC-1994
ATTORNATION NUMBER: BP 94104160.0
FILING DATE: 03-243-6313
TELEPHONE: 03-243-6313
TELEPHONE: 03-243-6313
TELEPHONE: 03-243-6313
TELEPHONE: 03-243-6310
INFORMATION FOR SEQ ID NO: 14:
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SEQUENCE CHARACTERISTICS:

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; OTHER INFORMATION: VL of scFv from mouse immunized with 2-phenyl-5-oxazolone US-09-726-219A-247
                                                                                                                                                                                                                                                                                                                                                                                                         FILE REFERENCE: 213839-00013
CURRENT APPLICATION NUMBER: US/09/726,219A
CURRENT FILING DATE: 2000-11-28
PRIOR APPLICATION NUMBER: GB 9015198.6
PRIOR FILING DATE: 1990-07-10
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR FILING DATE: 1990-10-19
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR FILING DATE: 1990-10-19
PRIOR APPLICATION NUMBER: GB 9024503.6
PRIOR APPLICATION NUMBER: GB 9024503.6
PRIOR FILING DATE: 1990-11-12
PRIOR FILING DATE: 1990-11-12
PRIOR FILING DATE: 1990-11-16
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  Query Match
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SOFTWARE: PatentIn version
SEQ ID NO 247
LENGTH: 110
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Patent No. 6806079
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Winter, Gregory
APPLICANT: Winter, Timothy
TITLE OF INVENTION: Methods for Producing Members of Specific Binding Pairs
FILE REFERENCE: 213839-00013
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APPLICANT: Cambridge Antibody Technology Limited
APPLICANT: Medical Research Council
                                                                                        TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 1991-07-10
                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: PCT/GB91/01134
                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 1991-05-15
                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US 07/971,857 FILING DATE: 1993-01-08
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Clackson, Timothy
Chiswell, David
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Griffiths, Andrew
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Pred. No. 4.9e-46;
  Score 541;
  DB 2;
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Length 110;
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CURRENT APPLICATION NUMBER: US/09/196,522

CURRENT FILING DATE: 1998-11-28

PRIOR APPLICATION NUMBER: GB 9015198.6

PRIOR FILING DATE: 1990-07-10

PRIOR APPLICATION NUMBER: GB 9022845.3

PRIOR FILING DATE: 1990-10-19

PRIOR APPLICATION NUMBER: GB 9022845.3

PRIOR APPLICATION NUMBER: GB 9022845.3

PRIOR APPLICATION NUMBER: GB 902885.3

PRIOR FILING DATE: 1990-10-19

PRIOR APPLICATION NUMBER: GB 9024503.6

PRIOR APPLICATION NUMBER: GB 9024744.9

PRIOR PRILING DATE: 1990-11-12

PRIOR APPLICATION NUMBER: GB 9104744.9

PRIOR FILING DATE: 1990-11-12

PRIOR APPLICATION NUMBER: GB 9104744.9

PRIOR FILING DATE: 1990-10-19
                                                                                                                                                                                                                                                                                                                                            SOFTWARE: Pa
SEQ ID NO 247
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                                                                                                                                                                     Query Match
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APPLICANT:
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APPLICANT: Bonert, Timothy
TITLE OF INVENTION: Methods for Producing Members of Specific Binding
FILE REFERENCE: 213839-00004
                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQ ID NOS: 272
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ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: VL of scFv from mouse immunized with 2-phenyl-5-oxazolone
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OR FILING DATE: 1991-05-15
OR APPLICATION NUMBER: PCT/GB91/01134
OR FILING DATE: 1991-07-10
OR APPLICATION NUMBER: US 07/971,857
OR FILING DATE: 1993-01-08
OR APPLICATION NUMBER: US 08/484,893
OR APPLICATION NUMBER: US 08/484,893
OR FILING DATE: 1995-06-07
                                                                                                                                                   Local
  61 ARFSGSGSGTSYSLTIGTMEAEDVATYYCQQGSSIPRTFGGGTKLEIKR 109
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                                                                                                                                                   Similarity
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                                                                                    DSELTQSPTTMAASPGEKITTTCSASSSISSNYLHWYQQRPGFSPKLLIYRTSNLASGVP
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Cambridge Antibody Technology Limited
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Griffiths, Andrew
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96.3%;
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Pred. No. 2.3e-45;
1; Mismatches 3
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1; Mismatches 3
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RESULT 4 US-09-726-219A-249

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APPLICANT: Winter, Gregory
APPLICANT: Bonert, Timothy
ITILE OF INVENTION: Methods for Producing Members of Specific Binding Pairs
FILE REFERENCE: 213839-00013
CURRENT APPLICATION NUMBER: US/09/726,219A
CURRENT FILING DATE: 2000-11-28
PRIOR APPLICATION NUMBER: GB 9015198.6
PRIOR APPLICATION NUMBER: GB 902845.3
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR FILING DATE: 1990-10-19
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR APPLICATION NUMBER: GB 9024503.6
PRIOR APPLICATION NUMBER: GB 9024503.6
PRIOR APPLICATION NUMBER: GB 904744.9
PRIOR FILING DATE: 1990-11-12
PRIOR FILING DATE: 1991-03-06
PRIOR APPLICATION NUMBER: GB 9104744.9
PRIOR FILING DATE: 1991-03-06
RESULT 5
US-09-196-522-249
; Sequence 249, Application US/09196522
; Patent No. 6916605
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SOFTWARE: PatentIn version 3.1
SEQ ID NO 249
LENGTH: 110
TYPE: PRT
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APPLICANT:
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PRIOR APPLICATION NUMBER: US 08/484,893
PRIOR FILING DATE: 1995-06-07
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PRIOR APPLICATION NUMBER: PCT/GB91/01134
PRIOR FILING DATE: 1991-07-10
PRIOR APPLICATION NUMBER: US 07/971,857
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                                                                                                                                                                                                                                                                                                                                          Local Similarity
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Griffiths, Andrew
Jackson, Ronald
                                                                                                                                      ARFSGSGSGTSYSLTIGTMEAEDVATYYCQQGSSIPFTFGGGTKLBIKR
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Sequence 245, Application US/09726219A Patent No. 6806079
GENERAL INFORMATION:

APPLICANT: Cambridge Antibody Technology
APPLICANT: Cambridge Antibody Technology Limited
APPLICANT: Medical Research Council
APPLICANT: McCafferty, John

APPLICANT:

Pope, Anthony Johnson, Kevin Hoogenboom, Hendricus

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US-09-726-219A-245
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Matches
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LENGTH: 110
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APPLICANT: Bonert, Timothy
TITLE OP INVENTION: Methods for Producing Members of Specific Binding Pairs
FILE REFERENCE: 213839-00004
CURRENT FILING DATE: 1998-11-28
PRIOR APPLICATION NUMBER: GB 9015198.6
PRIOR FILING DATE: 1990-07-10
PRIOR FILING DATE: 1990-07-10
PRIOR APPLICATION NUMBER: GB 902845.3
PRIOR PILING DATE: 1990-10-19
PRIOR PILING DATE: 1990-10-19
PRIOR PILING DATE: 1990-10-19
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NUMBER OF SEQ ID NOS: 272
SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: US 07/971,857
PRIOR FILING DATE: 1993-01-08
PRIOR APPLICATION NUMBER: US 08/484,893
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PRIOR FILING DATE: 1991-03-06
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FILING DATE: 1991-05-15
APPLICATION NUMBER: PCT/GB91/01134
FILING DATE: 1991-07-10
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                                                                                                              61 ARFSGSGSGTSYSLTIGTMEAEDVATYYCQQGSSIPRTFGGGTKLEIKR 109
                                                                                      61 ARPSGSGSGTSYSLTIGTMEAEDVATYYCQQGSSIPFTFGGGTKLBIKR
                                                                                                                                                                          1 DIELTŐS PTTMAAS PGEKITITCS AS SSIS SNYLHWY QOKPGPS PKLLIYRTSNLAS GVP
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Cambridge Antibody Technology Limited
Medical Research Council
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Griffiths, Andrew
Jackson, Ronald
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                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                95.98;
                                                                                                                                                                                                                                                            Score 540; DB 2;
Pred. No. 2.9e-45;
1; Mismatches 3
                                                                                                                                                                                                                                                                                                      Length 110;
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APPLICANT: APPLICANT: APPLICANT:

Chiswell, David

Timothy

James

PPLICANT:

Griffiths,

Andrew

Ronald

Kasper

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US-09-196-522-245
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                                                                                                                                                                                                                                                                                               Sequence 245, Application US/09196522 Patent No. 6916605
GENERAL INFORMATION:
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SOPTWARE: PatentIn version 3.1
SEQ ID NO 245
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Best Local Similarity
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CURRENT APPLICATION NUMBER: US/09/726,219A
CURRENT FILING DATE: 2000-11-28
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TITLE OF INVENTION: Methods for Producing Members of Specific Binding Pairs
FILE REFERENCE: 213839-00013
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APPLICANT: Cambridge Antibody Technology Limited
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PRIOR FILING DATE: 1990-07-10
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR FILING DATE: 1990-10-19
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ORGANISM: Artificial Sequence
PEATURE:
TYPE: PRT
ORGANISM: WI. of scFV
    PPLICANT:
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FILING DATE: 1993-01-08
APPLICATION NUMBER: US 08/484,893
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                                                                         Hoogenboom, Hendricus
Griffiths, Andrew
Jackson, Ronald
Holliger, Kasper
Marks, James
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                                  Chiswell, David
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Winter, Gregory
Bonert, Timothy
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95.4%;
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Pred. No. 5.7e-45;
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Best Local Similarity
Matches 104; Conserv
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CURRENT APPLICATION NUMBER: US/09/726,219A
CURRENT FILING DATE: 2000-11-28
PRIOR APPLICATION NUMBER: GB 9015198.6
PRIOR FILING DATE: 1990-07-10
PRIOR APPLICATION NUMBER: GB 9022845.3
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                                                                                                     APPLICANT: Winter, Gregory
APPLICANT: Bonart, Timothy
TITLE OF INVENTION: Methods for Producing Members of Specific Binding Pairs
FILE REFERENCE: 213839-00013
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CURRENT APPLICATION NUMBER: US/09/196,522
CURRENT FILING DATE: 1998-11-28
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FILING DATE: 1991-07-10
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Holliger, Kasper
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Hoogenboom, Hei
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McCafferty, John
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                                                                                                                                                                                                                                                                                                                                                       Anthony
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Pred. No. 5.7e-45;
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GB 9022845.3

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APPLICANT: Winter, Gregory
APPLICANT: Bonert, Timothy
TITLE OP INVENTION: Methods for Producing Members of Specific Binding Pair
FILE REFERENCE: 213839-00004
CURRENT FILING DATE: 1998-11-28
CURRENT FILING DATE: 1998-11-28
PRIOR APPLICATION NUMBER: GB 9015198.6
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR FILING DATE: 1990-10-19
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR APPLICATION NUMBER: GB 9024503.6
PRIOR APPLICATION NUMBER: GB 9024503.6
PRIOR APPLICATION NUMBER: GB 904503.6
PRIOR APPLICATION NUMBER: GB 904503.6
PRIOR FILING DATE: 1990-10-19
PRIOR APPLICATION NUMBER: GB 904503.6
PRIOR PRIOR DATE: 1991-03-06
PRIOR FILING DATE: 1991-03-06
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PRIOR PRIOR PRIOR NUMBER: GB 9104744.9
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US-09-196-522-258
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SEQ ID NO 258
LENGTH: 110
TYPE: PRT
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PRIOR FILING DATE: 1993-01-08
PRIOR APPLICATION NUMBER: US 08/484,893
PRIOR FILING DATE: 1995-06-07
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APPLICANT: Cambridge Antibody Technology
APPLICANT: Medical Research Council
APPLICANT: McGafferty, John
APPLICANT: Pope, Anthony
APPLICANT: Johnson, Kevin
APPLICANT: Johnson, Kevin
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Local Similarity 95.4%;
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APPLICATION NUMBER: GB 9104744.9
FILING DATE: 1991-03-06
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FILING DATE: 1991-05-15
APPLICATION NUMBER: PCT/GB91/01134
FILING DATE: 1991-07-10
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Griffiths, Andrew
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Marks, James
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1991-05-15
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Pred. No. 9e-45;
1; Mismatches
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; OTHER INFORMATION: US-09-196-522-258
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                                    PRIOR APPLICATION NUMBER: GB 9104744.9
PRIOR FILING DATE: 1991-03-06
PRIOR PELLOATION NUMBER: GB 9110549.4
PRIOR PELLOATION NUMBER: GB 9110549.4
PRIOR APPLICATION NUMBER: PCT/GB91/01134
PRIOR APPLICATION NUMBER: PCT/GB91/01134
PRIOR FILING DATE: 1991-07-10
PRIOR PELLOATION NUMBER: US 07/971,857
PRIOR PELLOATION NUMBER: US 07/971,857
PRIOR PELLOATION NUMBER: US 07/971,857
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LENGTH: 110
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Best Local Similarity
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PRIOR FILING DATE: 1995-06-07
NUMBER OF SEQ ID NOS: 272
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PRIOR FILING DATE: 1991-07-10
PRIOR APPLICATION NUMBER: US 07/971,857
PRIOR FILING DATE: 1993-01-08
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APPLICANT: Bonert, Timothy
TITLE OF INVENTION: Methods for Producing Members of Specific Binding Pairs
FILE REFERENCE: 213839-00013
CURRENT APPLICATION NUMBER: US/09/726,219A
CURRENT FILLING DATE: 2000-11-28
PRIOR APPLICATION NUMBER: GB 9015198.6
PRIOR PILLING DATE: 1990-07-10
PRIOR FILLING DATE: 1990-07-10
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PRIOR FILING DATE: 1990-10-19
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR FILING DATE: 1990-10-19
PRIOR APPLICATION NUMBER: GB 9024503.6
PRIOR FILING DATE: 1990-11-12
PRIOR FILING DATE: 1990-11-12
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TYPE: PRT
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APPLICANT: Cambridge Antibody Technology Limited
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                      APPLICATION NUMBER: US 08/484,893
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Griffiths, Andrew
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Pred. No. 9e-45;
1; Mismatches 4
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; OTHER INFORMATION: VL of scFv from mouse immunized with 2-phenyl-5-oxazolone US-09-726-219A-248
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US-09-726-219A-248
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                                                                                                    SOFTWARE: Pater
SOFTWARE: Pater
SEQ ID NO 248
SEQTH: 110
                                                                                                                                                                                               PRIOR APPLICATION NUMBER: GB 9110549.4
PRIOR FILING DATE: 1991-05-15
PRIOR PELICATION NUMBER: PCT/GB91/01134
PRIOR FILING DATE: 1991-07-10
PRIOR PELICATION NUMBER: US 07/971,857
PRIOR PELICATION NUMBER: US 07/971,857
PRIOR PELICATION NUMBER: US 08/484,893
PRIOR PELICATION NUMBER: US 08/484,893
PRIOR PILING DATE: 1995-06-07
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Best Local Similarity 94.5
Matches 103; Conservative
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TITLE OF INVENTION: Methods for Producing Members of Specific Binding Pairs

FILE REFERENCE: 213839-00013

CURRENT APPLICATION NUMBER: US/09/726,219A

CURRENT FILING DATS: 2000-11-28

PRIOR APPLICATION NUMBER: GB 9015198.6

PRIOR PILING DATS: 1990-07-10

PRIOR APPLICATION NUMBER: GB 9022845.3

PRIOR APPLICATION NUMBER: GB 9022845.3

PRIOR FILING DATS: 1990-10-19

PRIOR PILING DATS: 1990-10-19

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PRIOR FILING DATS: 1990-10-19
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ORGANISM: Artificial Sequence
FEATURE:
                                          ORGANISM: Artificial Sequence FEATURE:
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                                                                                                                                                                                               APPLICATION NUMBER: US 07/971,857
FILING DATE: 1993-01-08
APPLICATION NUMBER: US 08/484,893
FILING DATE: 1995-06-07
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Griffiths, Andrew
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Chiswell, David
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Pred. No. 1.8e-44;
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; OTHER INFORMATION: VL
US-09-196-522-238
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                                      Query Match
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APPLICANT: Bonert, Timothy
TITLE OF INVENTION: Methods for Producing Members of Specific Binding Pairs
FILE REFERENCE: 213839-00004
CURRENT APPLICATION NUMBER: US/99/196,522
CURRENT FILING DATE: 1998-11-28
PRIOR APPLICATION NUMBER: GB 9015198.6
PRIOR FILING DATE: 1990-07-10
PRIOR APPLICATION NUMBER: GB 902845.3
PRIOR PELICATION NUMBER: GB 9022845.3
PRIOR FILING DATE: 1990-10-19
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR FILING DATE: 1990-10-19
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                                                                                                                                                                                                                                       NUMBER OF SEQ ID NOS: 272
                                                                                                                           TYPE: PRT
ORGANISM: Artificial Sequence
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                                                                                                                    FEATURE:
                                                                                                                                                                                ENGTH: 110
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FILING DATE: 1991-05-15
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                        Similarity
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Griffiths, Andrew
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94.5%;
ilarity 94.5%;
Conservative
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                                                                                                  of scFv from mouse immunized with 2-phenyl-5-oxazolone
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  Score 532; DB 2;
Pred. No. 1.8e-44;
2; Mismatches 4
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Pred. No. 1.8e-44;
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                                         Length 110
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DSELTQSPTTMAASPGEKITTTCSASSSISSNYLHWYQQRPGFSPKLLIYRTSNLASGVP

DIBLTQSPTTMAASPGEKITITCSASSSISSNYLHWYQQKPGFSPKLLIYRTSNLASGVP

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RESULT 13
US-09-196-522-248
                      RESULT 14
US-09-486-814A-2
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PRIOR FILING DATE: 1991-03-06
PRIOR APPLICATION NUMBER: GB 9110549.4
PRIOR FILING DATE: 1991-05-15
PRIOR APPLICATION NUMBER: PCT/GB91/01134
PRIOR FILING DATE: 1991-07-10
PRIOR FILING DATE: 1993-01-08
PRIOR FILING DATE: 1993-01-08
PRIOR FILING DATE: 1993-01-08
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Patent No. 6916605
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SOFTWARE: PatentIn version 3.1
SEQ ID NO 248
LENGTH: 110
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Sequence 2, Application US/09486814A
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APPLICANT: Bonert, Timothy
TITLE OF INVENTION: Methods for Producing Members of Specific Binding Pairs
FILE REFERENCE: 213839-00004
CURRENT APPLICATION NUMBER: US/09/196,522
CURRENT FILING DATE: 1998-11-28
PRIOR APPLICATION NUMBER: GB 9015198.6
PRIOR PILING DATE: 1990-07-10
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR FILING DATE: 1990-10-19
PRIOR APPLICATION NUMBER: GB 9022845.3
                                                                                                                                                                                                                                                                                       Matches 103;
                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
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OTHER INFORMATION: VL of scFv from mouse immunized with 2-phenyl-5-oxazolone
-09-196-522-248
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APPLICANT: Cambridge Antibody Technology Limited
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PRIOR APPLICATION NUMBER: GB 9024503.6
PRIOR FILING DATE: 1990-11-12
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                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US 08/484,893
                                                                                                             61
                                                                                                                                                    61 ARFSGSGSGTSYSLTIGTMEAEDVATYYCQQGSSIPRTFGGGTKLEIKR 109
                                                                                                                                                                                                                       DSELTQSPTTMAASPGEKITTTCSASSSISSNYLHWYQQRPGFSPKLLIYRTSNLASGVP
                                                                                                                                                                                                  DIELTOS PTTWAAS PGEKITITCS ASSSISSINLHWYQQKPGFS PKLLIYRTSNLASGVP
                                                                                                             ARPSGSGSGTSYSLTIGTMEAEDVATYYCQQGSGIPYTFGGGTKLBIKR 109
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McCafferty, John
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Holliger, Kasper
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US-09-726-219A-250
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PRIOR APPLICATION NUMBER: GB 9015198.6
PRIOR FILING DATE: 1990-07-10
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR FILING DATE: 1990-10-19
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR FILING DATE: 1990-10-19
PRIOR FILING DATE: 1990-10-19
PRIOR PPLICATION NUMBER: GB 9024503.6
PRIOR APPLICATION NUMBER: GB 9024503.6
PRIOR APPLICATION NUMBER: GB 9024503.6
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GENERAL INFORMATION:
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Best Local Similarity 93.7%;
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APPLICANT: YAMAMOTO, Masato
APPLICANT: HAYASHI, No. 656259910
APPLICANT: YAMAMOTO, Hiroko
APPLICANT: TOHDOH, Naoki
TITLE OF INVENTION: SINGEE-CHAIN ANTIBODY AGAINST HEPATITIS B VIRUS CORE
TITLE OF INVENTION: PROTEIN, GENE THEREOF, AND THERAPEUTIC AGENT FOR
TITLE OF INVENTION: HEPATITIS B CONTAINING THESE
FILE REFERENCE: 0020-4682P
TITLE OF INVENTION: HEPATITIS B CONTAINING THESE
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APPLICANT:
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APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
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                                                                                                                                                                        APPLICANT: Winter, Gregory
APPLICANT: Bonert, Timothy
TITLE OF INVENTION: Methods for Producing Members of Specific Binding Pairs
FILE REFERENCE: 213839-00013
CURRENT APPLICATION NUMBER: US/09/726,219A
CURRENT FILING DATE: 2000-11-28
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ORGANISM: Mus sp., strain: Balb/c, tissue: spleen
FEATURE:
NAME/KEY: PEPTIDE
LOCATION: (1)...(145)
OTHER INFORMATION: Identification Method: P
                                                                                                                                                                                                                                                                                                         APPLICANT:
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LOCATION: (177)..(279)
OTHER INFORMATION: Identification Method:
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Cambridge Antibody Technology Limited
Medical Research Council
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Clackson, Timothy
Chiswell, David
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Griffiths, Andrew
Jackson, Ronald
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Pred. No. 8.7e-44;
2; Mismatches 3
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PRIOR APPLICATION NUMBER: GB 9104744.9

PRIOR FILING DATE: 1991-03-06

PRIOR APPLICATION NUMBER: GB 9110549.4

PRIOR APPLICATION NUMBER: GB 9110549.4

PRIOR PILING DATE: 1991-05-15

PRIOR APPLICATION NUMBER: PCT/GB91/01134

PRIOR PILING DATE: 1991-07-10

PRIOR APPLICATION NUMBER: US 07/971,857

PRIOR PILING DATE: 1993-01-08

PRIOR PILING DATE: 1993-06-07

NUMBER: US 08/484,893

PRIOR PILING DATE: 1995-06-07

NUMBER OF SEQ ID NOS: 272

SOFTWARE: Patentin version 3.1

SEQ ID NO 250

LENGTH: 110

TYPE: PRT

ORGANISM: Artificial Sequence

PEATURE:

OTHER INFORMATION: VL of scFv from mouse immunized with 2-phenyl-5-oxazolone

US-09-726-219A-250
Search completed: November 21, 2005, 12:07:37 Job time: 42.1245 secs
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                                                                              61 ARFSGSGSGTSYSLTIGTMEAEDVATYYCQQGSSIPRTFGGGTKLEIKR 109
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Maximum DB
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of hits satisfying chosen parameters:
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1: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*

2: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

3: /cgn2_6/ptodata/1/pubpaa/US09_PUBCOMB.pep:*

4: /cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*

5: /cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.pep:*
Listing first 45 summaries
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US-09-144-886-92
US-10-632-706-89
US-10-803-622-247
US-10-803-653-247
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US-10-803-653-245
US-10-803-622-245
US-10-803-622-248
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US-10-803-622-248
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US-10-803-622-250
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4 US-10-803-693-255     Sequence 255       4 US-10-803-693-256     Sequence 256       4 US-10-803-622-257     Sequence 257       4 US-10-803-653-257     Sequence 257       5 US-10-803-653-257     Sequence 257       6 US-10-803-622-237     Sequence 237       7 US-10-803-622-237     Sequence 237       8 US-10-803-653-237     Sequence 240       9 US-10-803-653-240     Sequence 240       9 US-10-803-653-240     Sequence 240       9 US-10-803-653-254     Sequence 254       9 US-10-803-652-254     Sequence 254       9 US-10-803-653-254     Sequence 254       9 US-10-461-80-653-25     Sequence 254       9 US-10-461-80-006-20     Sequence 20       9 US-10-461-805-12     Sequence 21       10 US-10-461-805-12     Sequence 21       10 US-10-461-805-90     Sequence 20       10 US-10-461-806-90     Seq	78.5	78.7	78.7	78.9	78.9	78.9	78.9	78.9	79.4	79.4	79.9	79.9	79.9	79.9	80.3	80.3	80.8	80.8	
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-10-803-693-255 -10-803-693-256 Sequence 256 Sequence 257 -10-803-623-257 Sequence 257 -10-803-623-257 Sequence 257 Sequence 257 Sequence 257 Sequence 257 Sequence 257 Sequence 257 Sequence 237 Sequence 237 Sequence 237 Sequence 237 Sequence 240 Sequence 240 Sequence 240 Sequence 240 Sequence 257 -10-803-653-254 Sequence 254 -10-803-653-254 Sequence 254 -10-803-653-254 Sequence 254 -10-803-653-254 Sequence 254 -10-803-653-254 Sequence 254 -10-803-653-254 Sequence 254 -10-461-878-12 Sequence 20, -10-461-885-12 Sequence 20, -10-461-885-12 Sequence 30, -09-144-886-90 Sequence 90,	ω	v	4	4	4.	4	4	4.	4	4	4	4	4	w	4	4	4	4	
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## ALIGNMENTS

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RESULT 1
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US-10-632-706-89
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APPLICANT: Marks, James D
APPLICANT: Amersdorfer, Peter
APPLICANT: Amersdorfer, Peter
TITLE OF INVENTION: Therapeutic Monoclonal Antibodies That Neutralize
TITLE OF INVENTION: Botulinum Neurotoxins
FILE REFERENCE: 2500.117USO
CURRENT APPLICATION NUMBER: US/09/144,886
CURRENT APPLICATION NUMBER: US/09/144,886
CURRENT FILING DATE: 1998-08-31
NUMBER OF SEQ ID NOS: 98
SOPTWARE: Patentin Ver. 2.0
SEQ ID NO SEQ ID NO
                                                                                                                                               Sequence 89, Application US/10632706
publication No. US20040175385A1
GENERAL INFORMATION:
APPLICANT: MARKS, JAMES D.
APPLICANT: AMERGDORFER, PETER
APPLICANT: AMERGDORFER, PETER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 92, Appl Patent No. US2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
APPLICANT: AMERSDORFER, PETER
TITLE OF INVENTION: THERAPEUTIC MONOCLONAL ANTIBODIES THAT NEUTRALIZE BOTULINUM
TITLE OF INVENTION: NEUROTOXINS
FILE REFERENCE: 4077-895120US
CURRENT APPLICATION NUMBER: US/10/632,706
CURRENT FILING DATE: 2003-08-01
PRIOR APPLICATION NUMBER: US 60/400,721
PRIOR APPLICATION NUMBER: US 60/400,721
PRIOR FILING DATE: 2002-08-01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 109
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OTHER INFORMATION: Description of Artificial Sequence: BONT/A clone OTHER INFORMATION: 183 region VL epitope 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
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S20020155114A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
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PRIOR APPLICATION NUMBER: US 09/144,806
PRIOR FILING DATE: 1998-08-31
NUMBER OF SEQ ID NOS: 278
SOFTWARE: Patentin version 3.2
SEQ ID NO 89
LENGTH: 109
TYPE: PRT
ORGANISM: Artificial
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                                                               SOFTWARE: Patentin
SEQ ID NO 247
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Publication No. US20040157214A1
                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Winter, Gregory
APPLICANT: Bonert, Timothy
TITLE OF INVENTION: Methods for Producing Members of Specific Binding
FILE REFERENCE: 11839-00013
CURRENT APPLICATION NUMBER: US/10/803,622
CURRENT APPLICATION NUMBER: GB 9015198.6
PRIOR APPLICATION NUMBER: GB 9015198.6
PRIOR PILING DATE: 1990-07-10
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR PILING DATE: 1990-10-19
PRIOR PILING DATE: 1990-10-19
PRIOR PILING DATE: 1990-10-19
PRIOR PILING DATE: 1990-10-19
PRIOR PILING DATE: 1990-10-10-19
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR PILING DATE: 1990-10-10-10
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                                                                                                          PRIOR FILING DATE: 1995-0
NUMBER OF SEQ ID NOS: 272
                                                                                                                                                       PRIOR FILING DATE: 1991-07-10

PRIOR APPLICATION NUMBER: US 07/971,857

PRIOR FILING DATE: 1993-01-08

PRIOR APPLICATION NUMBER: US 08/484,893
                                                                                                                                                                                                                                      PRIOR FILING DATE: 1991-05-15
PRIOR APPLICATION NUMBER: PCT/GB91/01134
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Cambridge Antibody Technology
APPLICANT: Cambridge Antibody Technology Limited
APPLICANT: Medical Research Council
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Local Similarity 100.0%;
                                                                                                                                                     APPLICATION NUMBER: US 08/484,893
                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: GB 9024503.6 FILING DATE: 1990-11-12 APPLICATION NUMBER: GB 9104744.9
                                                                                                                                                                                                                                                                                                            FILING DATE: 1991-03-06
                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: GB 9110549.4
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Holliger, Kasper
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Griffiths, Andrew
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                                                                                        version 3.1
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; Pred. No. 5.7e-42;
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APPLICANT: Winter, Gregory
APPLICANT: Bonert, Timothy
ITITLE OF INVENTION: Methods for Producing Members of Specific Binding Pair
FILE REFERENCE: 13839-00013
CURRENT APPLICATION NUMBER: US/10/803,653
CURRENT FILING DATE: 2004-03-18
PRIOR APPLICATION NUMBER: GB 9015198.6
PRIOR FILING DATE: 1990-07-10
PRIOR PILING DATE: 1990-10-19
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR FILING DATE: 1990-10-19
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR FILING DATE: 1990-10-19
PRIOR FILING DATE: 1990-10-19
PRIOR FILING DATE: 1990-11-12
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PRIOR FILING DATE: 1990-11-12
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US-10-803-653-247
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; OTHER INFORMATION: VL of scFv from mouse immunized with 2-phenyl-5-oxazolone
US-10-803-622-247
                                                                                                           SOFTWARE: Par
SEQ ID NO 247
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT:
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                                                                                                                                                   PRIOR FILING DATE: 1995-0.
NUMBER OF SEQ ID NOS: 272
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches 105; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: GB 9110549.4
                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: GB 9104744.9 PRIOR FILING DATE: 1991-03-06
        OTHER INFORMATION: VL
                               FEATURE:
                                              TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                          ENGTH:
                                                                                                                                                                                                                           RILING DATE: 1991-05-15
APPLICATION NUMBER: PCT/GB91/01134
FILING DATE: 1991-07-10
APPLICATION NUMBER: US 07/971,857
                                                                                                                                                                                         FILING DATE: 1993-01-08
APPLICATION NUMBER: US 08/484,893
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Griffiths, Andrew
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p. US20040157215A1
                                                                                                                                                                      1995-06-07
of scFv from mouse immunized with 2-phenyl-5-oxazolone
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Pred. No. 5e-40;
1; Mismatches
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Query Match 96.1 Best Local Similarity 96.3 Matches 105; Conservative

96.1%; Score 541; DB 4 96.3%; Pred. No. 5e-40; cive 1; Mismatches

DB 4, Length 110

Indels

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Gaps

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APPLICANT: Winter, Gregory
APPLICANT: Bonert, Timothy
IIILS OP INVENTION, Methods for Producing Members of Specific Binding Pair
FILE REFERENCE: 13839-00013
CURRENT FILING DATE: 2004-03-18
CURRENT FILING DATE: 2004-03-18
PRIOR APPLICATION NUMBER: GB 9015198.6
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR FILING DATE: 1990-10-19
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR FILING DATE: 1990-10-19
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR FILING DATE: 1990-10-19
PRIOR FILING DATE: 1990-10-19
PRIOR APPLICATION NUMBER: GB 9024503.6
PRIOR FILING DATE: 1990-11-12
PRIOR APPLICATION NUMBER: GB 9104744.9
PRIOR APPLICATION NUMBER: GB 9104744.9
PRIOR APPLICATION NUMBER: GB 9104744.9
PRIOR APPLICATION NUMBER: GB 9104744.9
PRIOR APPLICATION NUMBER: GB 910549.4
PRIOR APPLICATION NUMBER: GB 9110549.4
PRIOR APPLICATION NUMBER: GB 910549.4
PRIOR APPLICATION NUMBER: GB 910549.4
PRIOR PILING DATE: 1991-07-10
PRIOR APPLICATION NUMBER: US 07/971,857
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US-10-803-622-249
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SEQ ID NO 249
LENGTH: 110
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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GENERAL INFORMATION
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Matches
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APPLICANT: Cambridge Antibody Technology Limited
APPLICANT: Medical Research Council
APPLICANT: McCafferty, John
APPLICANT: Pope, Anthony
APPLICANT: Hoogenboom, Hendricus
                                                                                                                                                                                                                                                     OTHER INFORMATION: VL of scFv from mouse immunized with 2-phenyl-5-oxazolone 10-803-622-249
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                                                                                                                                                                                   Local
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105; Conserv
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                                                                                                     DSELTQSPTTMAASPGEKITTTCSASSSISSNYLHWYQQRPGFSPKLLIYRTSNLASGVP
  ARFSGSGSGTSYSLTIGTMEAEDVATYYCQQGSSIPFTFGGGTKLEIKR
                              ARFSGSGSGTSYSLTIGTMKARDVATTYYCQQGSSIPRTFGGGTKLEIKR 109
                                                                                 DIELTQSPTTMAASPGEKITITCSASSSISSNYLHWYQQKPGFSPKLLIYRTSNLASGVP
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Griffiths, Andrew
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b. US20040157214A1
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Pred. No. 6.1e-40
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RESULT 6
US-10-803-653-249
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; OTHER INFORMATION: VL of scFv US-10-803-653-249
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CURRENT FILING DATE: 2004-03-18
PRIOR APPLICATION NUMBER: GB 9015198.6
PRIOR APPLICATION NUMBER: GB 9015198.6
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR PILING DATE: 1990-10-19
PRIOR PILING DATE: 1990-10-19
PRIOR PILING DATE: 1990-11-12
PRIOR APPLICATION NUMBER: GB 9024503.6
PRIOR APPLICATION NUMBER: GB 9104744.9
PRIOR APPLICATION NUMBER: GB 9104744.9
PRIOR APPLICATION NUMBER: GB 9104744.9
PRIOR APPLICATION NUMBER: GB 9110549.4
PRIOR PILING DATE: 1991-03-06
PRIOR PILING DATE: 1991-03-06
PRIOR PILING DATE: 1991-05-15
PRIOR PILING DATE: 1991-05-15
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SEQ ID NO 249
LENGTH: 110
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Publication No. US20040157215A1
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Best Local Similarity
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APPLICANT: Cambridge Antibody Technology Limited
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PRIOR FILING DATE: 1993-01-08
PRIOR APPLICATION NUMBER: US 08/484,893
PRIOR FILING DATE: 1995-06-07
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APPLICANT: Bonert, Timothy
TITLE OF INVENTION: Methods for Producing Members of Specific Binding Pairs
FILE REFERENCE: 13839-00013
                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQ ID NOS: 272
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: PCT/GB91/01134 PRIOR FILING DATE: 1991-07-10
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                                     61 ARFSGSGSGTSYSLTIGTMEAEDVATYYCQdGSSIPRTFGGGTKLEIKR
                                                                                                                             1 DSELTQSPTTMAASPGEKITTTCSASSSIS9NYLHWYQQRPGFSPKLLIYRTSNLASGVP
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Griffiths, Andrew
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                                                                                                                                                                           Conservative
                                                                                                                                                                                                  95.98;
                                                                                                                                                                           1; Mismatches
                                                                                                                                                                                                  Score 540; DB 4;
Pred. No. 6.1e-40;
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US-10-803-622-245

Sequence 245, Application US/10803622 Publication No. US20040157214A1 GENERAL INFORMATION:

APPLICANT: Cambridge Antibody Technology AppLICANT: Cambridge Antibody Technology Limited

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US-10-803-653-245

1 Sequence 245, Application US/10803653

1 Publication No. US20040157215A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; CTHER: INFORMATION: VL of scFv from mouse immunized with 2-phenyl-5-oxazolone
US-10-803-622-245
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PRIOR FILING DATE: 1990-07-10
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR FILING DATE: 1990-10-19
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR FILING DATE: 1990-10-19
PRIOR APPLICATION NUMBER: GB 90245.3
PRIOR APPLICATION NUMBER: GB 90245.3.6
PRIOR APPLICATION NUMBER: GB 9024503.6
PRIOR APPLICATION NUMBER: GB 9024503.6
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NUMBER OF SEQ ID NOS: 272
SOFTWARE: PatentIn version 3.1
SEQ ID NO 245
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
                                                                                                           APPLICANT: Cambridge Antibody Technology
APPLICANT: Cambridge Antibody Technology Limited
APPLICANT: Medical Research Council
APPLICANT: McCafferty, John
APPLICANT: Pope, Anthony
APPLICANT
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PRIOR FILING DATE: 1991-05-15
PRIOR APPLICATION NUMBER: PCT/GB91/01134
PRIOR FILING DATE: 1991-07-10
PRIOR APPLICATION NUMBER: US 07/971,857
PRIOR FILING DATE: 1993-01-08
PRIOR APPLICATION NUMBER: US 08/484,893
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APPLICANT: Bonert, Timothy
TITLE OF INVENTION: Methods for Producing Members of Specific Binding
FILE REFERENCE: 13839-00013
CURRENT APPLICATION NUMBER: US/10/803,622
CURRENT FILING DATE: 2004-03-18
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PRIOR FILING DATE: 1991-03-06
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              Hoogenboom, Hendricus
Griffiths, Andrew
Jackson, Ronald
                                                                                  Johnson, Kevin
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McCafferty, John
Pope, Anthony
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PRIOR FILING DATE: 1990-07-10
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR FILING DATE: 1990-10-19
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR FILING DATE: 1990-10-19
PRIOR PILING DATE: 1990-10-19
PRIOR APPLICATION NUMBER: GB 9024503.6
PRIOR FILING DATE: 1990-11-12
PRIOR FILING DATE: 1990-11-12
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LENGTH: 110
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Best Local
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                                                                                                         APPLICANT:
PPLICANT: Winter, Gregory

APPLICANT: Bonert, Timothy

TITLE OF INVENTION: Methods for Producing Members of Specific Binding Pairs

TLE REFERENCE: 13839-00013
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: US 08/484,893
PRIOR FILING DATE: 1995-06-07
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APPLICANT: Bonert, Timothy
TITLE OF INVENTION: Methods for Producing Members of Specific Binding Pairs
FILE REFERENCE: 13839-00013
CURRENT APPLICATION NUMBER: US/10/803,653
CURRENT FILING DATE: 2004-03-18
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ORGANISM: Artificial Sequence
FEATURE:
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APPLICATION NUMBER: PCT/GB91/01134
FILING DATE: 1991-07-10
APPLICATION NUMBER: US 07/971,857
FILING DATE: 1993-01-08
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Cambridge Antibody Technology Limited
Medical Research Council
McCafferty, John
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                                                                               Clackson, Timothy Chiswell, David
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Griffiths, Andrew
                                                                                                                                          Jackson, Ronald
Holliger, Kasper
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o. US20040157214A1
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Chiswell, David
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CURRENT APPLICATION NUMBER: US/10/803,622

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PRIOR APPLICATION NUMBER: GB 9015198.6
PRIOR FILING DATE: 1990-07-10
PRIOR PRIOR PAPELICATION NUMBER: GB 9022845.3
PRIOR FILING DATE: 1990-10-19
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR FILING DATE: 1990-10-19
PRIOR APPLICATION NUMBER: GB 9024503.6
PRIOR APPLICATION NUMBER: GB 9104744.9
PRIOR FILING DATE: 1991-03-06
PRIOR PRIOR DATE: 1991-03-06
PRIOR APPLICATION NUMBER: GB 9110549.4
PRIOR FILING DATE: 1991-05-16
PRIOR APPLICATION NUMBER: BD 9110549.4
PRIOR FILING DATE: 1991-07-10
PRIOR APPLICATION NUMBER: US 07/971,857
PRIOR APPLICATION NUMBER: US 08/484,893
PRIOR PILING DATE: 1993-01-08
PRIOR APPLICATION NUMBER: US 08/484,893
PRIOR FILING DATE: 1995-06-07
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US-10-803-653-258
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SEQ ID NO 258
LENGTH: 110
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Best Local Similarity 95.4%;
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                                                                 APPLICANT: Winter, Gregory
APPLICANT: Bonett, Timothy
TITLE OF INVENTION: Methods for Producing Members of Specific Binding Pairs
FILE REFERENCE: 13839-00013
CURRENT APPLICATION NUMBER: US/10/803,653
CURRENT FILING DATE: 2004-03-18
PRIOR APPLICATION NUMBER: GB 9015198.6
PRIOR APPLICATION NUMBER: GB 9015198.6
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR FILING DATE: 1990-07-10
PRIOR FILING DATE: 1990-10-19
PRIOR FILING DATE: 1990-10-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Cambridge Antibody Technology
APPLICANT: Cambridge Antibody Technology Limited
APPLICANT: Medical Research Council
APPLICANT: McCafferty, John
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                          APPLICATION NUMBER: GB 9022845.3 FILING DATE: 1990-10-19
APPLICATION NUMBER: GB 9024503.6
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Griffiths, Andrew
Jackson, Ronald
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Marks, James
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o. US20040157215A1
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Pred. No. 1.7e-39;
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RESULT 12 US-10-803-622-238

Sequence 238, Application US/10803622

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OTHER INFORMATION: VL of scPv from mouse immunized with 2-phenyl-5-oxazolone US-10-803-653-258
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Patent No. US20020155114A1
GENERAL INFORMATION:
APPLICANT: Marks, James D
APPLICANT: Marks, James D
TITLE OF INVENTION: Therapeutic Monoclonal Antibodies That Neutralize
TITLE OF INVENTION: Therapeutic Monoclonal Antibodies That Neutralize
TITLE OF INVENTION: WOUNDER: WS/09/144,886
FILE REFERENCE: 2500.117USO
CURRENT APPLICATION NUMBER: US/09/144,886
CURRENT APPLICATION NUMBER: US/09/144,886
CURRENT PILING DATE: 1998-08-31
NUMBER OF SEQ ID NOS: 98
SOPTWARE: PatentIn Ver. 2.0
SEQ ID NO 98
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SOFTWARE: PatentIn version 3.1
SEQ ID NO 258
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Best Local &
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TYPE: PRT
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                                                                                                                                                                                                                                               FEATURE: OTHER INFORMATION: Description of Artificial Sequence: BoNT/A clone OTHER INFORMATION: 2E8 region VL epitope 4
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                                                                                                                                                  Local Similarity 94.
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APPLICATION NUMBER: PCT/GB91/01134
FILING DATE: 1991-07-10
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                    61 ARPSGSGSGTSYSLTIGTMEAEDVATYYCQQGSSIPRTFGGGTKLEIKR 109
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ARFSGSGSGTSYSLTIGAMEAEDVATYYCOOGSSIPYTFGGGTKLEIKR
                                                                         DIELTQSPTTMAASPGEKITITCSASSSIGSNYLHWYQQKPGFSPKLLIYRTSNLASGVP
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Pred. No. 1.7e-39;
1; Mismatches 4
                                                                                                                                                  Score 532; DB 3;
Pred. No. 3.1e-39;
1; Mismatches 5
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Publication No. US20040157214A1

Cambridge Antibody Technology

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SEQ ID NO 238
LENGTH: 110
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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US-10-803-622-248
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US-10-803-622-238
                                                                                                            Sequence 248, Application US/10803622 Publication No. US20040157214A1 GENERAL INFORMATION:
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Best Local Similarity
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PRIOR APPLICATION NUMBER: PCT/GB91/01134
PRIOR FILING DATE: 1991-07-10
PRIOR APPLICATION NUMBER: US 07/971,857
PRIOR FILING DATE: 1993-01-08
PRIOR FILING DATE: 1993-01-08
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TITLE OF INVENTION: Methods for Producing Members of Specific Binding Pairs
FILE REFERENCE: 13839-00013
CURRENT APPLICATION NUMBER: US/10/803,622
CURRENT FILING DATE: 2004-03-18
PRIOR APPLICATION NUMBER: GB 9015198.6
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR PILING DATE: 1990-07-10
PRIOR PPLICATION NUMBER: GB 9022845.3
PRIOR PILING DATE: 1990-10-19
PRIOR PILING DATE: 1990-10-19
PRIOR FILING DATE: 1990-10-19
PRIOR FILING DATE: 1990-10-19
PRIOR FILING DATE: 1990-10-19
PRIOR PILING DATE: 1990-10-19
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                     APPLICANT: Cambridge Antibody Technology Limited APPLICANT: Cambridge Antibody Technology Limited APPLICANT: Medical Research Council APPLICANT: McCafferty, John
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PRIOR APPLICATION NUMBER: GB 9104744.9
PRIOR FILING DATE: 1991-03-06
PRIOR APPLICATION NUMBER: GB 9110549.4
PRIOR FILING DATE: 1991-05-15
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NUMBER OF SEQ ID NOS: 272
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Griffiths, Andrew
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Marks, James
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Pope, Anthony
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Timothy
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APPLICANT: Griffiths, Andrew
APPLICANT: Jackson, Ronald
APPLICANT: Holliger, Kasper
APPLICANT: Holliger, Kasper
APPLICANT: Clackson, Timothy
APPLICANT: Clackson, Timothy
APPLICANT: Clackson, Timothy
APPLICANT: Clackson, Timothy
TITLE Clackson, Timothy
TITLE BONER, Gregory
APPLICANT: BONER, Timothy
TITLE REFERENCE: 13839-00013
CURRENT FILING DATE: 13939-0013
CURRENT FILING DATE: 2004-03-18
PRIOR APPLICATION NUMBER: GB 9015198.6
PRIOR APPLICATION NUMBER: GB 902845.3
PRIOR FILING DATE: 1990-10-19
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR FILING DATE: 1990-10-19
PRIOR PILING DATE: 1990-10-19
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR APPLICATION NUMBER: GB 9022845.3
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PRIOR FILING DATE: 1990-11-12
PRIOR PPLICATION NUMBER: GB 9104744.9
PRIOR PPLICATION NUMBER: GB 9104744.9
PRIOR APPLICATION NUMBER: GB 9110549.4
PRIOR FILING DATE: 1991-05-15
PRIOR APPLICATION NUMBER: PCT/GB91/01134
PRIOR PILING DATE: 1991-07-10
PRIOR APPLICATION NUMBER: US 07/971,857
PRIOR APPLICATION NUMBER: US 07/971,857
PRIOR APPLICATION NUMBER: US 08/484,893
PRIOR PILING DATE: 1993-01-08
PRIOR APPLICATION NUMBER: US 08/484,893
PRIOR FILING DATE: 1995-06-07
                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                    Sequence 238, Appropriate Publication No.
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Best Local Similarity 94.5%;
Matches 103; Conservative
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SOFTWARE: PatentIn versio
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     APPLICANT
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ORGANISM: Artificial Sequence
FEATURE:
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Jackson, Ronald
Holliger, Kasper
Marks, James
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Hoogenboom, Hendricus
Griffiths, Andrew
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o. US20040157215A1
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Chackson, Timothy Chiswell, David

Gregory

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PRIOR FILING DATE: 1991-07-10
PRIOR APPLICATION NUMBER: US 07/971,857
PRIOR FILING DATE: 1993-01-08
PRIOR APPLICATION NUMBER: US 08/484,893
PRIOR FILING DATE: 1995-06-07
NUMBER OF SEQ ID NOS: 272
SOFTWARE: Patentin version 3.1
SEQ ID NO 238
LENGTH: 110
TYPE: PRT
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; OTHER INFORMATION: VL of scFv from mouse immunized with 2-phenyl-5-oxazolone
US-10-803-653-238
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PRIOR FILING DATE: 1991-03-06
PRIOR APPLICATION NUMBER: GB 9110549.4
PRIOR FILING DATE: 1991-05-15
PRIOR APPLICATION NUMBER: PCT/GB91/01134
PRIOR APPLICATION NUMBER: US 07/971,857
PRIOR FILING DATE: 1993-01-08
PRIOR APPLICATION NUMBER: US 08/484,893
PRIOR APPLICATION NUMBER: US 08/484,893
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US-10-803-653-248
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Best Local Similarity
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TITLE OF INVENTION: Methods for Producing Members of Specific Binding

FILE REFERENCE: 13839-00013

CURRENT APPLICATION NUMBER: US/10/803,653

CURRENT FILING DATE: 2004-03-18

PRIOR APPLICATION NUMBER: GB 9015198.6

PRIOR FILING DATE: 1990-07-10

PRIOR PRIOR DATE: 1990-10-19

PRIOR FILING DATE: 1990-10-19

PRIOR FILING DATE: 1990-10-19

PRIOR APPLICATION NUMBER: GB 9022845.3

PRIOR FILING DATE: 1990-10-19

PRIOR APPLICATION NUMBER: GB 9024503.6

PRIOR APPLICATION NUMBER: GB 9024503.6

PRIOR FILING DATE: 1990-11-12

PRIOR FILING DATE: 1990-11-12

PRIOR FILING DATE: 1990-11-12

PRIOR FILING DATE: 1990-11-12
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APPLICANT:
APPLICANT: Winter, Gregory
APPLICANT: Bonert, Timothy
TITLE OP INVENTION: Methods for Producing Members of Specific Binding
FILE REFERENCE: 13839-00013
CURRENT APPLICATION NUMBER: US/10/803,653
CURRENT FILING DATE: 2004-03-18
PRIOR APPLICATION NUMBER: GB 9015198.6
PRIOR APPLICATION NUMBER: GB 9015198.6
PRIOR APPLICATION NUMBER: GB 9022845.3
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Marks, James
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Griffiths, Andrew
Jackson, Ronald
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o. US20040157215A1
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Pred. No. 3.1e-39;
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, OTHER INFORMATION: VL of scFv from mouse immunized with 2-phenyl-5-oxazolone US-10-803-653-248
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NUMBER OF SEQ ID NOS: 272
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PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR FILING DATE: 1990-10-19
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ORGANISM: Artificial Sequence
PEATURE:
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                                                                                                                                                                    Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: GB 9024503.6 FILING DATE: 1990-11-12 APPLICATION NUMBER: GB 9104744.9 FILING DATE: 1991-03-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: GB 9110549.4
FILING DATE: 1991-05-15
APPLICATION NUMBER: PCT/GB91/01134
FILING DATE: 1991-07-10
                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US 07/971,857 FILING DATE: 1993-01-08
                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US 08/484,893
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                         61 ARPSGSGSGTSYSLTIGTMEAEDVATYYCQÓGSSIPRTFGGGTKLEIKR
61 ARFSGSGSGTSYSITIGTMEAEDVATYYCQQGSGIPYTFGGGTKLEIKR 109
                                                                                              1 DSELTQSPTTMAASPGEKITTTCSASSSISSNYLHWYQQRPGFSPKLLIYRTSNLASGVP
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Pred. No. 3.1e-39;
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Copyright (c) 1993 - 2005 Compugen Ltd.
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US-11-174-186-7
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US-11-174-186-16
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Sequence 1, Appli
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Sequence 19, Appli
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Similarity 75.2%; pred. No. 3.4e-19; 9; CORBETVALIVE 7; Mismatches 17; LTQSPTTMAASPGEKITTTCSASSISSNYLHWYQOREG	186-1 186-1 170, Application US/11174186 11 Application US/11174186 11 INFORMATION: 10 Stephen 10 INT: Gillies, Stephen 10 INT: Gillies, Stephen 10 INT: Lo, Kin-Ming 10 INT: Lo, Kin-Ming 10 INT: Lo, Kin-Ming 10 INT: Lo, Kin-Ming 10 INT: Lo, Kin-Ming 10 INTERIOR IN	26 367.5 65.3 543 1 US-10-495-644-3 Sequence 3, Appli 27 355.5 63.1 129 1 US-10-721-743-19 Sequence 19, Appli 28 354.5 63.0 111 7 US-11-77-978-1 Sequence 1, Appli 39.4 5.5 62.6 131 1 US-10-721-763-27 Sequence 27, Appli 39.4 5.5 61.5 126 1 US-10-839-799-65 Sequence 27, Appli 39.4 5.5 61.4 126 1 US-10-839-799-65 Sequence 47, Appli 34.5 61.4 126 1 US-10-839-799-85 Sequence 65, Appli 36 345.5 61.4 126 1 US-10-839-799-85 Sequence 65, Appli 39.4 5.5 61.4 126 1 US-10-839-799-85 Sequence 65, Appli 34.5 61.4 126 1 US-10-839-799-85 Sequence 65, Appli 39.4 5.5 61.4 126 1 US-10-819-799-85 Sequence 65, Appli 34.5 61.0 127 1 US-11-779-78-4 Sequence 68, Appli 39.4 5.5 61.4 126 1 US-10-71-184-31 Sequence 31, Appli 39.4 5.5 61.0 127 1 US-11-779-78-4 Sequence 4, Appli 39.4 5.5 61.0 236 7 US-11-779-78-4 Sequence 34, Appli 39.4 5.5 61.0 236 7 US-11-144-288-48 Sequence 34, Appli 39.4 5.5 61.0 259 1 US-10-512-184-34 Sequence 34, Appli 39.4 5.5 61.0 259 1 US-10-512-184-34 Sequence 29, Appli 39.4 5.5 61.0 256 1 US-10-512-184-71 Sequence 29, Appli 39.4 5.5 61.0 371 1 US-10-512-184-71 Sequence 29, Appli 39.4 5.5 61.0 371 1 US-10-512-184-71 Sequence 34, Appli 39.4 5.5 61.0 371 1 US-10-512-184-71 Sequence 37, Appli 39.4 5.5 61.0 626 1 US-10-839-799-43 Sequence 39, Appli 39.5 61.0 526 1 US-10-512-184-71 Sequence 39, Appli 39.5 61.0 526 1 US-10-839-799-53 Sequence 39, Appli 39.5 61.0 526 1 US-10-839-799-53 Sequence 39, Appli 39.5 61.0 526 1 US-10-839-799-53 Sequence 39, Appli 39.5 61.0 526 1 US-10-839-799-53 Sequence 39, Appli 39.5 61.0 526 1 US-10-839-799-53 Sequence 39, Appli 39.5 61.0 526 1 US-10-839-799-53 Sequence 39, Appli 39.5 61.0 526 1 US-10-839-799-53 Sequence 39, Appli 39.5 61.0 526 1 US-10-839-799-53 Sequence 39, Appli 39.5 61.0 526 1 US-10-839-799-53 Sequence 59, Appli 39.
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CURRENT APPLICATION NUMBER: US/11/174,186

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PRIOR APPLICATION NUMBER: US 60/288,564
PRIOR FILING DATE: 2001-05-03
NUMBER OF SEQ ID NOS: 42
SOFTWARE: Patentin version 3.3
SEQ ID NO 9
LENGTH: 106
TYPE: PRT
; Sequence 7, Application US/11174186; Publication No. US20050244418A1; GENERAL INFORMATION:
; APPLICANT: Gillies, Stephen; APPLICANT: Lo, Kin-Ming; APPLICANT: Qian, Xiugi; TITLE OF INVENTION: Recombinant Tumor Sp.; FILE REFERENCE: LEX-019
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US-11-174-186-7
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APPLICANT: Gillies, Stephen
APPLICANT: II, Kin-Ming
APPLICANT: Qian, Xiugi
TITLE OF INVENTION: Recombinant Tumor Specific Antibody And Use Thereof
FILE REFERENCE: LEX-019
CURRENT APPLICATION NUMBER: US/11/174,186
CURRENT FILING DATE: 2005-07-01
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PRIOR APPLICATION NUMBER: US 60/288,564
PRIOR FILING DATE: 2001-05-03
NUMBER OF SEQ ID NOS: 42
SOFTWARE: PatentIn version 3.3
SEQ ID NO 42
LENGTH: 213
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ORGANISM: Artificial
PRATURE:
OTHER INFORMATION: light chain
3-11-174-186-42
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Best Local &
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OTHER INFORMATION: VK8 light chain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Artificial FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       / Match 70.7%; Score 398; DB 7; Length 213; Local Similarity 73.6%; Pred. No. 4.1e-19; ndels 78; Conservative 9; Mismatches 17; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                        4 LTQSPTTMAASPGEKITTTCSASSSISSNYLHWYQQRPGFSPKLLIYRTSNLASGVPARF 63
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                                                                                                                                                                                                                                                                                                                                           LTQSPATLSLSPGERVTLTCSASSSVS--YMLWYQQXPGSSPXPWIFDTSNLASGFPARF 61
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                                                                                                                                                                                                                                                                                                                                                                                                                                  9; Mismatches
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                          Specific Antibody And Use Thereof
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RESULT 6 US-11-174-186-8

; Sequence 8, Application US/11174186

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GENERAL INFORMATION:

APPLICANT: Adolf, Guenther

APPLICANT: Baum, Anke

APPLICANT: Heider, Karl-Heinz

ITILE OF INVENTION: Compositions and Methods for Treating Cancer using

ITILE OF INVENTION: Cytotoxic CD44 Antibody Immunoconjugates and

ITILE OF INVENTION: Chemotherapeutic Agents

FILE REPERENCE: 1/1383

CURRENT APPLICATION NUMBER: US/11/172,320

CURRENT APPLICATION NUMBER: US/10/645,215

PRIOR APPLICATION NUMBER: US/00/645,215

PRIOR PILING DATE: August 26, 2002

NUMBER OF SEQ ID NOS: 9

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US-11-172-320-4
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CURRENT FILING DATE: 2005-07-01
PRIOR APPLICATION NUMBER: US 60/288,564
PRIOR FILING DATE: 2001-05-03
NUMBER OF SEQ ID NOS: 42
                                                                                                                                                                                                                                                Best Local Similarity
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TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: Vk6 light chain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Artificial
                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: Humanised Murine Antibody BIWA 4 Light Chain
                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: PRT
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                                                                                                                                                          4 LTOSPTTMAASPGEKITTTCSASSSISSNYLHWYQQRPGFSPKLLIYRTSNLASGVPARF 63
                                                                                                                                                                                                                   75; Conservative
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                                        SGSGSGTSYSLTIGTMEASDVATYYCQQGSSIPRTFGGGTKLEIKR 109
                                                                                                         LTQSPATLSLSPGERATLSCSASSSI--NYIYWYQQKPGQAPRLLIYLTSNLASGVPARF 61
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                                                                                                                                                                                                             69.4%; Score 391; DB 7; Length 213
70.8%; Pred. No. 1.1e-18;
tive 14; Mismatches 15; Indels
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Best Local Similarity
Thes 77; Conserv
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SEQ ID NO 8
LENGTH: 106
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 36, Application US/10512184 Publication No. US20050244901A1
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PRIOR APPLICATION NUMBER: US 60/288,564
PRIOR FILING DATE: 2001-05-03
NUMBER OF SEQ ID NOS: 42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Lo, Kin-Ming
APPLICANT: Olan, Xiugi
TITLE OJAN, Xiugi
TITLE OLIVENTION. Steephinant Tumor Specific Antibody And Use Thereof
FILE REFERENCE: LEX-019
CURRENT APPLICATION NUMBER: US/11/174,186
CURRENT FILING DATE: 2005-07-01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT FILING DATE: 2004-10-22 NUMBER OF SEQ ID NOS: 72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Fraunhofer Gesellschaft zur F*rderung der angewandten Forschung e.V. TITLE OF INVENTION: Antibodies, recombinant antibodies, recombinant TITLE OF INVENTION: antibody fragments and fusions mediated plant disease TITLE OF INVENTION: resistance against fungi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILE REFERENCE: 3581.01US01
CURRENT APPLICATION NUMBER: US/10/512,184
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Gillies,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FEATURE: OTHER INFORMATION: VK7 light chain
                                                                                                                                                                                                                                                                        OTHER INFORMATION: Description of Artificial Sequence: scFv SS2 with OTHER INFORMATION: specificity against Sclerotinia sclerotiorum; OTHER INFORMATION: originates from Mus musculus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Artificial
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                                                                                                                                                                                                                                                                                                                                             FEATURE:
                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 248
                                                                                                                                                                                        Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION:
                                                                                  141 DIVLEOSPVIMSASPGEKVTMTCSASSSV--NYIYWYQWKSGTSPKRWIFDTSKLASGVP
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                          61 ARFSGSGSGTSYSLTIGTMEAEDVATYYCQQGSSIPRTFGGGTKLEIKR 109
                                                                                                                          1 DSELTQSPTTMAASPGEKITTTCSASSSISSNYLHWYQQRPGFSPKLLIYRTSNLASGVP 60
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LTQSPAIMSASPGERVTMTCSASSSVS--YMLWYQQKPGSSPKDWIFDTSNLASGFPARF
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  VRFSGSGSGTSFSLTISSMEAEDIATYYCOOWSSPPLTFGAGTKLELKR 247
                                                                                                                                                                      Conservative
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70.6%;
                                                                                                                                                                      11; Mismatches
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Pred. No. 7.4e-19;
                                                                                                                                                                                          Score 390; DB 1;
Pred. No. 1.4e-18;
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RESULT 8 US-10-512-184-50

RESULT 10

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US-11-174-186-15
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Publication No. US20050244901A1
GENERAL INFORMATION:
APPLICANT: Fraunhofer Gesellschaft zur F"rderung der angewandten Forschung TITLE OF INVENTION: Antibodies, recombinant antibodies, recombinant TITLE OF INVENTION: antibody fragments and fusions mediated plant disease TITLE OF INVENTION: resistance against fungi
FILE REFERENCE: 3581.01US01
CURRENT FILING ADTE: 2004-10-22
NUMBER OF SEQ ID NOS: 72
SOFTMARE: Batentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                 FILE REFERENCE: LEX-019
CURRENT APPLICATION NUMBER: US/11/174,186
CURRENT FILING DATE: 2005-07-01
PRIOR APPLICATION NUMBER: US 60/288,564
PRIOR FILING DATE: 2001-05-03
NUMBER OF SEQ ID NOS: 42
SOFTWARE: PATENTIN VERSION 3.3
SEQ ID NO 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 15, Application US/11174186 Publication No. US20050244418A1 GENERAL INFORMATION:
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LENGTH: 61
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                                                                                                                                                                                 Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: Recombinant Tumor Specific Antibody And Use
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Artificial Sequence FEATURE: FEATURE: OTHER INFORMATION: Description of Artificial Sequence: fusion OTHER INFORMATION: comprising the leader peptide - chitinase OTHER INFORMATION: - scFv SS2 - cmyc/His6.
                                                                                                                                                                                                                                                                                                     LENGTH: 106
TYPE: PRT
ORGANISM: Artificial
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62
                                      64 SGSGSGTSYSLTIGTMEAEDVATYYCQQGSSIPRTFGGGTKLBIK 108
                                                                                                                     4 LTOSPTTMAASPGEKITTTCSASSSISSNYLHWYQQRPGFSPKLLIYRTSNLASGVPARF
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                                                                             Qian, Xiugi
  SGSGSGTSYTLTINSLEAEDAATYYCHQRSGYPYTFGGGTKVEIK 106
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                                                                                                                                                                                   68.7%; Score 387; DB 7; 71.4%; Pred. No. 1.1e-18;
                                                                                                                                                                11; Mismatches
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US-11-174-186-16

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GENERAL INFORMATION: OSCIOLARIANAL GENERAL INFORMATION: APPLICANT: Adolf, Guenther APPLICANT: Heider, Karl-Heinz APPLICANT: Heider, Karl-Heinz TITLE OF INVENTION: Compositions and Methods for Treating Cancer using TITLE OF INVENTION: Cytotoxic CD44 Antibody Immunoconjugates and TITLE OF INVENTION: Chemotherapeutic Agents FILE REFERENCE: 1/1383
CURRENT PILING DATE: 2005-06-30
PRIOR APPLICATION NUMBER: US/10/645,215
PRIOR APPLICATION NUMBER: US/10/645,215
PRIOR APPLICATION NUMBER: 20 018 686.2
PRIOR APPLICATION NUMBER: 21, 2002
PRIOR APPLICATION NUMBER: US 60/405,956
PRIOR FILING DATE: August 21, 2002
PRIOR APPLICATION NUMBER: US 60/405,956
PRIOR FILING DATE: August 26, 2002
PRIOR FILING DATE: August 26, 2002
PRIOR FILING DATE: August 26, 2002
PRIOR FILING DATE: August 26, 2002
PRIOR FILING DATE: August 26, 2002
PRIOR FILING DATE: August 26, 2002
PRIOR FILING DATE: August 26, 2002
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US-11-172-320-8
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; OTHER INFORMATION: KS VK mouse US-11-174-186-16
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CURRENT APPLICATION NUMBER: US/11/174,186
CURRENT FILING DATE: 2005-07-01
PRIOR APPLICATION NUMBER: US 60/288,564
PRIOR FILING DATE: 2001-05-03
NUMBER OF SEQ ID NOS: 42
SOFTWARE: Patentin version 3.3
SEQ ID NO 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 8, Application US/11172320 Publication No. US20050244413A1
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Publication No. US20050244418A1
                                                                                                               Matches
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Best Local |
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                                                                                                                                                                                                                                                                             TYPE: PRT
ORGANISM: Artificial Sequence
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ORGANISM: Artificial
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                                                                                                            Similarity 73; Conserv
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74; Conser
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                                                                                                            67.7%; Score 381; DB 7; ilarity 68.9%; Pred. No. 4.2e-18; Conservative 15; Mismatches 16
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                                                                                                                                                                DB 7; Length 213;
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; ORGANISM: Homo sapiens
US-10-648-816-5
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-648-816-1
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US-10-648-816-1
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US-10-648-816-5
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APPLICANT: Van Bruggen, Nicholas
APPLICANT: Perrara, Napoleone
TITLE OF INVENTION: Vascular Endothelial Cell Growth Factor Antagonists
TITLE OF INVENTION: Vascular Endothelial Cell Growth Factor Antagonists
TITLE OF INVENTION: Vascular Endothelial Cell Growth Factor Antagonists
TILLE REFERENCE: P1717D1
CURRENT APPLICATION NUMBER: US/10/648,816
CURRENT FILING DATE: 2003-08-26
PRIOR APPLICATION NUMBER: US/09/718,694
PRIOR FILING DATE: 2000-11-21
PRIOR APPLICATION NUMBER: US 09/218,481
PRIOR APPLICATION NUMBER: US 09/218,481
PRIOR FILING DATE: 1998-12-22
NUMBER OF SEQ ID NOS: 16
SEQ ID NO 5
                                                                                                                                                                                                                                                           SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 1, Application US/10648816
Publication No. US20050244405A1
GENERAL INFORMATION:
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Best Local :
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                                                                                                            Best Local
                                                                                                                           Query Match
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APPLICANT: Ferrara, Napoleone
TITLE OF INVENTION: Vascular Endothelial Cell Growth Factor Antagonists
TITLE OF INVENTION: and Uses Thereof
FILE REFERENCE: P1717D1
                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/10/648,816
CURRENT FILING DATE: 2003-08-26
PRIOR PPLICATION NUMBER: US/09/718,694
PRIOR FILING DATE: 2000-11-21
PRIOR APPLICATION NUMBER: US 09/218,481
PRIOR FILING DATE: 1998-12-22
PRIOR FILING DATE: 1998-12-22
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ch 67.4%; Score 379.5; DB 1; 1 Similarity 65.1%; Pred. No. 3.2e-18; 71; Conservative 22; Mismatches 15;
                                                                                                          Similarity
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                                                                                      Conservative
                                                                                                          67.1%; Score 377.5; DB 1; 64.2%; Pred. No. 4.2e-18;
                                                                                        Mismatches
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                                                                                                                             Length 110;
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61 ARPSGSGSGTSYSLTIGTMEAEDVATYYCQQGSSIPRTFGGGTXLEIKR 109
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Sequence 14, Application US/11174186

Publication No. US20050244418A1

GENERAL INFORMATION:
APPLICANT: Gillies, Stephen'
APPLICANT: Lo, Kin-Ming
APPLICANT: Qian, Xiugi
ITILE OF INVENTION: Recombinant Tumor Specific Antibody And Use Thereof
FILE REPERBNCE: LEX-019
CURRENT APPLICATION NUMBER: US/11/174,186
CURRENT APPLICATION NUMBER: US/11/174,186
CURRENT APPLICATION NUMBER: US/11/174,186
CURRENT APPLICATION NUMBER: US/07-01
PRIOR APPLICATION NUMBER: US/07-03
NUMBER OF SEQ ID NOS: 42
SOPTWARE: Patentin version 3.3
SEQ ID NO 14
LENGTH: 106
TYPE: PRT
ORGANISM: Artificial
PEATURE:
COMMENT OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TABLE OF TAB
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US-11-174-186-10
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US-11-174-186-14
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CURRENT FILING DATE: 2005-07-01
PRIOR APPLICATION NUMBER: US 60/288,564
PRIOR FILING DATE: 2001-05-03
NUMBER OF SEQ ID NOS: 42
SOFTWARE: Patentin version 3.3
SEQ ID NO 10
LENGTH: 106
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Publication No. US20050244418A1
GENERAL INFORMATION:
APPLICANT: Gillies, Stephen
APPLICANT: Lo, Kin-Ming
APPLICANT: Qian, Xiugi
TITLE OF INVENTION: Recombinant Tumor Specific Antibody And Use Thereof
FILE REFERENCE: LEX-019
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Matches
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                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                OTHER INFORMATION: KS de-immunized VK4
11-174-186-14
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                                             LTQSPTTMAASPGEKITTTCSASSSISSNYLHWYQQRPGFSPXLLIYRTSNLASGVPARF 63
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    LTQSPASLAVSPGQRATITCSASSSVS--YMLWYQQKPGQPPKPWIFDTSNLASGFPARF 61
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                      64 SGSGSGTSYSLTIGTMEAEDVATYYCQQGSSIPRTFGGGTKLBIK 108
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Search completed: November 21, 2005, 12:33:51
Job time : 3.09634 secs

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Minimum DB seq length: 0
Maximum DB seq length: 200000000
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## SUMMARIBS

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ADRIBU antibacterial; antibody; botulinum neurotoxin type A; BoNT/A; BoNT/A neutralising epitope; anti-botulinum neurotoxin antibody; toxin neutralisation; botulinum neurotoxin poisoning; mouse; heavy chain variable region; complementarity determining region; CDR3. ADR38724 standard; peptide; 11 AA. 01-AUG-2003; 2003US-00632706 09-SEP-2004. US2004175385-A1. Mus sp. Mouse heavy 02-DEC-2004 ADR38724; chain variable region CDR3 seqid 126. (first entry)

31-AUG-1998; 98US-00144886. 01-AUG-2002; 2002US-0400721P. (REGC ) UNIV CALIFORNIA.

New isolated antibody that neutralizes botulinum neurotoxin type A, useful for diagnosing botulism or for treating pathologies associated with botulinum neurotoxin poisoning.

WPI; 2004-652009/63.

Marks JD,

Amersdorfer P;

Example 3; SEQ ID NO 126; 110pp; English.

The invention describes an isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a specific clone where (I) binds to and neutralises botulinum neurotoxin type A (BoNT/A). An isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a clone chosen from clone 25, C35, C35, 1C6, 3D12, B4, 1F3, huC25, A11, A22, WR1(V), WR1(T), 3-1, 3-8, 3-10 and ING1, where (I) binds to and neutralizes botulinum neurotoxin type A (BoNT/A). Also described are: a polypeptide (II) comprising BoNT/A neutralising epitope having an epitope that is

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RESULT 2
ADR38725
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CC bountinum neurotoxin antibodies, where each antibody is specific for a 
CC different epitope of a botulinum neurotoxin and the combination of 
CC antibodies shows greater toxin neutralisation than the single antibodies 
CC in surplus. The following are disclosed: a pharmaceutical composition 
CC comprising (I); and a kit comprising (I). (I) is useful for neutralising 
CC BNNTA antibody and for neutralising a botulinum neurotoxin which 
CC involves contacting neurotoxin with (I) in surplus, where each of (I) is 
CC combination of antibodies shows greater toxin neutralisation than the 
CC combination of antibodies shows greater toxin neutralisation than the 
CC combination of antibodies shows greater toxin neutralisation than the 
CC combination pathologies associated with botulinum neurotoxin 
CC poisoning. (I) exhibits specificity and affinity towards BoNT/A. (I) 
CC enables rapid detection or diagnosis of botulism. This is the amino acid 
CC sequence of mouse heavy chain variable region complementarity determining 
CC region 3 (CDR3) from anti-botulinum neurotoxin antibodies.
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Best Local Similarity
Matches 11; Conserv
  The invention describes an isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a specific clone where (I) binds to and neutralises botulinum neurotoxin type A (BoNT/A). An isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a clone chosen from clone $25, C25, C39, 1C6, 3D12, B4, 1F3, huC25, Ar1, Ar2, WR1(V), WR1(T), 3-1,
                                                                                                                                                                                                         New isolated antibody that neutralizes botulinum neurotoxin type A, useful for diagnosing botulism or for treating pathologies associated
                                                                                                                                                                                        with botulinum neurotoxin
                                                                                                                                                                                                                                                                                                             Marks JD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BONT/A neutralising epitope; anti-botulinum neurotoxin antibody; toxin neutralisation; botulinum neurotoxin poisoning; mouse; heavy chain variable region; complementarity determining region; CDR3.
                                                                                                                                              Example 3; SEQ ID NO 127; 110pp; English.
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01-AUG-2002; 2002US-0400721P.
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                                                                                                                                                                                                                                                                                        07-JAN-2005; 2005WO-US000370
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human monocional anti-MAdCAM antibody related protein
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                                                                                                                                                                                                                 (PFIZ ) PFIZER INC
                                                                                                                                                                                                                                                                                                                                                                                                                                  lmmunosuppressive;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   region 3 (CDR3) from anti-botulinum neurotoxin antibodies.
                                                                                                                                                                             ABGENIX INC. PFIZER LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LATYYYFGLDV 11
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                                                                                                                                         Molloy E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first
                                                                                                                                                                                                                                                                                                                                                                                                                                      antibody.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%; Score 60; 100.0%; Pred. No.
                                                                                                                                           Kellermann
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                                                                                                                                           Green
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                                                                                                                                           Haak-Frendscho
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New antibody to Mucosal Adressin Cell Adhesion Molecule, useful for diagnosing and treating an inflammatory disease, e.g. inflammatory disease, ulcerative colitis, gastritis, insulin-dependent diabetes graft versus host disease.

bowel

2005-554958/56.

Example 5, Fig 1, 167pp; English

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The invention relates to a human monoclonal antibody or its antigence adhesion molecule (MadCAM). The invention also relates to a hybridoma cell line that produces the human monoclonal antibody, a pharmaceutical composition comprising an amount of the monoclonal antibody or its composition comprising an amount of the monoclonal antibody or its produces the monoclonal antibody or its produces the monoclonal antibody or its produces the monoclonal antibody or its produces the monoclonal antibody or its produces the monoclonal antibody or its produces the monoclonal antibody or its produces the monoclonal antibody or its produces the monoclonal antibody or its produces the monoclonal antibody or its produces the monoclonal antibody or its produces the monoclonal antibody or its produces the monoclonal antibody or its private the heavy or its private the monoclonal comprising the monoclonal antibody or its private the monoclonal comprising the expression control sequence operably linked to the nucleic acid molecule, where the vector optionally comprises an expression control sequence operably linked to the nucleic acid molecule, where the vector of the nucleic acid molecule, where the vector of the nucleic acid molecule, or its antigen-binding portion that specifically binds of visual states and its antigen-binding portion that specifically binds of indeating an antibody or its antigen-binding portion that specifically binds to MAdCAM, and inhibits binding to emphasion the specifically binds to MAdCAM and inhibits binding to emphasion the specifically binds to MAdCAM and inhibits binding to emphasion the specifically binds to MAdCAM and inhibits binding to emphasion the specifically binds to MAdCAM and inhibits binding to emphasion the specifically binds to MAdCAM and inhibits binding to emphasion and infiltration into tissues, a method of inhibiting alphaseta? MAdCAM mad inhibits binding to the worth of disease, e.g. characterized by circulating soluble human MAdCAM and inhibits binding to the monoclonal methods a
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                                                                                                                                                                                                               B-cell; malignant; immunoglobulin; immunoglobulin variable region; Ig variable region; glycosylation site; lymphoma; B cell receptor; cytostatic; gene therapy; glycosylation inhibitor; non-Hodgkin's lymphoma.
                                                                                                                                                                              Synthetic
                                                                                                                                                                                                                                                                                                                                                            15-JAN-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                       ADD28082 standard; protein; 114 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          anti-MAdCAM antibody related protein of the invention.
  07-MAR-2002; 2002GB-00005395
                                       24-FEB-2003; 2003WO-GB000783
                                                                                                                  WO2003074059-A2
                                                                                                                                                                                                                                                                                                                   Lymphoma related immunoglobulin variable region.
                                                                              12-SEP-2003
                                                                                                                                                         Homo sapiens,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 LATYYYFGLDV 11
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2.5;
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The present invention describes a method for classifying a B-cell as CC malignant or normal comprising: (a) isolating a sequence representing an CC immunoglobulin (Ig) variable region from the B cell; (b) detecting the CC presence of a glycosylation site; and (d) classifying the cell as CC malignant or normal on the basis of the presence or absence of a CC glycosylation site. Also described: (1) treating a patient suffering from CC or at risk of having lymphoma; (2) screening for substances capable of inhibiting glycosylation of the Ig variable region of the B cell receptor CC; and (3) screening for substances (S) dapable of inhibiting the CC interaction between lectins of the type found in the germinal centre and CC meticant found on the surface of Ig of lymphoma cells. (8) has CC cytostatic activity, and can be used in gene therapy, and as a CC glycosylation inhibitor. The method is useful in classifying a B-cell as CC medicament for treating non-Hodgkin's lymphoma. The present sequence CC represents an Ig variable region sequence which is used in the
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Matches
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                11-FEB-2002;
29-APR-2002;
27-SEP-2002;
25-NOV-2002;
                                                                                                                                                                                                                                                              human heterodimeric antibody; human; antibody; binding affinity; protective antigen; Bacillus anthracis; antibrax infection; cell receptor; edema factor; lethal factor; virucide; antibacterial; immunotherapy; anti-toxin; anti-infective; anthrax; botulinum; smallpox; Venezuelan equine encephalomyelitis virus; VEEV; West Nile virus; WNV.
                                                                                                                                                                                                                                                                                                                                                                Human heterodimeric antibody heavy chair variable region SEQ ID NO:101.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 114 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                     ADD28323;
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                                                                                                   11-FEB-2003; 2003WO-US004206.
                                                                                                                                      18-SEP-2003
                                                                                                                                                                                                         Homo
                                                                                                                                                                                                                                                            Venezuelan equine
                                                                                                                                                                                                       sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             106 TYYYYGMDV 114
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3 TYYYFGLDV 11
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7; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Stevenson F;
                ; 2002US-0356086P.
; 2002US-0376408P.
; 2002US-0414053P.
; 2002US-0428807P.
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                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                            encephalomyelitis virus;
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Pred. No.
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(ALEX-) ALEXION PHARM

INC

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Matches 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention describes a human heterodimeric antibody (I) (fragment) having a binding affinity of at least 1x10-8 M to the protective antigen of Bacillus anthracis or a molecule involved in anthrax infection that blocks binding of the antigen or molecule to cell receptors, edema factor and lethal factor. (I) has virucide and antibacterial activities, and can be used in immunotherapy. The antibodies (I) are useful as anti-toxins or anti-infectives with respect to infective agents, such as anthrax, botulinum, smallpox, Venezuelan equine encephalomyelitis virus (VEEV), or West Nile virus (WNV). The present sequence represents a human heterodimeric antibody heavy chain variable region amino acid sequence, which is used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New human heterodimeric antibodies or their antibody fragments, useful as anti-toxins or anti-infectives with respect to infective agents, e.g. anthrax, botulinum, smallpox, Venezuelan equine encephalomyelitis or West
                                                                                                                                                                                                                     11-FEB-2002; 2002US-0356086P.
29-APR-2002; 2002US-0376408P.
27-SEP-2002; 2002US-0414063P.
25-NOV-2002; 2002US-0428807P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               human heterodimeric antibody; human; antibody; binding affinity;
protective antigen; Bacillus anthracis; anthrax infection; cell receptor;
edema factor; lethal factor; virucide; antibacterial; immunotherapy;
anti-toxin; anti-infective; anthrax; botulinum; smallpox;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 129
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                                                    WPI; 2003-722327/68
                                                                                                         Bowdish KS,
                                                                                                                                                                                                                                                                                                                                                                 11-FEB-2003; 2003WO-US004206
                                                                                                                                                                                                                                                                                                                                                                                                                      18-SEP-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic.
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                                                                                                                                                                   (ALEX-)
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7, Conserve
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                                                                                                                                                                ALEXION PHARM INC
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                                                                                                            Wild MA
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77.8%;
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Pred. No. 3.9;
2; Mismatches
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Matches 7
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comprises administering an antibody or antibody fragment that binds to a molecule involved in anthrax infection and which has the ability to block the binding of the molecule to at least one of a cell receptor, PA63, PA63 heptamer, PA83, edema factor and lethal factor. A claimed method for determining exposure to B. anthracis comprises assaying a sample for the presence of a molecule selected from cell receptors, PA63, PA63 heptamer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New human heterodimeric antibodies or their antibody fragments, useful as anti-toxins or anti-infectives with respect to infective agents, e.g. anthrax, botulinum, smallpox, Venezuelan equine encephalomyelitis or West
                                                                                                                                                                                                                                                                                                   Treating an animal with anthrax infection by administering the protective antigen of Bacillus anthracis, and blocking cell receptors, edema factor or lethal factor.
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                                                                                                                                                                                    A claimed method for treating an animal infected with Bacillus anthracis
                                                                                                                                                                                                                                              Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Bowdish
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Pred. No.
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Novel neuropilin binder which is scFv, antibody fragment or bioconjugate, that modulates neuropilin-1 function or inhibits NP-1 dependent angiogenesis of endothelial cells and/or invasion of tumor cells useful for treating cancer.

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15-JAN-2003; 2003EP-00000615
    WPI; 2004-507700/48
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                                                                                Bustace BK,
                                                                                Zehetmeier C, Lain B,
Knauer R, Jensen KH;
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                                                                                                                                Torella C,
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Niewoehner J;

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cc pa83, edema factor or lethal factor with an antibody that has binding cc affinity for the molecule, where the presence of elevated levels of the antibody correlates with the presence of a disease associated with B. cc anthracis. Alternatively, the method involves assaying for the presence of factor or lethal factor with a secondary antibody having binding affinity cf for the antibody, where the presence of elevated levels of the secondary critical variable region selected from a group binding affinity can light chain waitable region selected from a group of sequences ADV868127, a light chain kappa region selected from a group of sequences ADV868127, a light chain kappa region selected from a group or sequences compositions of the present inventional region selected from a group of sequences ADV868135 and a light chain lambda region selected from a group or sequences compositions of the present invention are also useful for producing antitoxing or anti-infectives to infective agents such as anthrax, botulinum, smallpox, Venezuelan equine encephalomyelitis and West Nile virus. The present sequence is that of the heavy chain variable region of a human compositions and bone marrow samples of domors who had been vaccinated sequence analysis was performed on positive resounders. Neutralization of anthrax toxin activity by purified Pabs was demonstrated.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 134 AA;
Cytostatic; metastasis inhibitor; neuropilin binder; NPB; scFv; single chain antibody; neuropilin-1; NP-1; angiogenesis; tumour
                                                                 NPB polypeptide scFv27, seq id
                                                                                                               07-OCT-2004
                                                                                                                                                            ADR28082
                                                                                                                                                                                                   ADR28082 standard; protein; 244 AA
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77.8%;
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    tumour; cancer.
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determining region

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Location/Qualifiers 87. .103

WO2004056874-A2

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                                                      Pullen N,
                                                                                                                                          (PFIZ) PFIZER INC.
(ABGE-) ABGENIX INC.
(PFIZ) PFIZER LTD.
                                                                                                                                                                                                                                                                                                                                       09-JAN-2004; 2004US-0535490P
                                                                                                                                                                                                                                                                                                                                                                                                                                     07-JAN-2005; 2005WO-US000370.
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                                                      Molloy E,
                                                      Kellermann
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                                                           Green
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                                                           Haak-Frendscho
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CC polypeptide, antibody, scFv, antibody fragment or bioconjugate, that CC modulates neuropilin-1 (NP-1) function for inhibits NP-1 dependent CC angiogenesis of endothelial cells and/or invasion of tumour cells, CC whereby the NPB binds to NP-1 and modulates NP-1 function. Further CC disclosed is an ex vivo method of determining the dependency of the CC invasiveness of a naturally occurring invasive cancer cell on the CC invasiveness of NP-1 mediated invasion and/or adhesion and an inhibitor of CC metastasis of NP-1 mediated invasion and/or adhesion and an inhibitor of CC is useful for detecting NP-1 expression, modulation of NP-1 function, CC adhesion of cells, preferably tumour cells. The NPB of the invention of antifacture of medicament for the treatment or prevention of NP-dependent and non-physiological blood vessel growth, particularly correlated with a tumour. It is also useful for treatment or prevention of cells, The current sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 3; SEQ ID NO 29; 120pp; English
                                      represents a single chain antibody neuropilin binder (NPB) polypeptide
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Matches Query Match Best Local S Sequence 244 AA; 95 Similarity
7; Conserve TYYYYGMDV 103 TYYYFGLDV 11 Conservative 78.3**%**; 77.8**%**; Score 47; DB Pred. No. 7.7; 2; Mismatches 2 8 0 Length 244 0 Gaps

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06-OCT-2005 AEB45891 standard; protein; 468 (first entry)

Human monoclonal anti-MAdCAM antibody #37.

WPI; 2005-554958/56. N-PSDB; AEB45890.

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AEB45853
ID AEB45853
XX AEB4585
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KW infize
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cell line that produces the human monoclonal antibody, a pharmaceutical composition comprising an amount of the monoclonal antibody or its composition comprising an amount of the monoclonal antibody or its cracking inflammatory disease in a subject, an isolated cell line that produces the monoclonal antibody or its antigen-binding portion or the cheavy chain or light chain of the antibody or of its portion, an isolated culcilated acid molecule comprising a nucleotide sequence encoding the heavy chain or its antigen-binding portion or the light chain or its antigen-binding portion or the light chain or its antigen-binding portion of an antibody described above, a vector comprising the vector optionally comprises an expression control sequence operably linked to the nucleic acid molecule, a host cell comprising the vector optionally comprises an expression that specifically binds MACAM, a method of isolating an antibody or its antigen-binding portion that specifically binds to MACAM, a method of its antigen-binding comprising the vector or the nucleic acid molecule, a method of treating a subject in need of a human antibody or its antigen-binding comprising to a subject in need of a human antibody or its antigen-binding to alphabeta? a method of inhibiting alphabeta? MACAM, a method of inhibiting madcham-mediated leukocyte composition of inhibiting alphabeta? MACAM-dependent cellular adhesion, composition and into tissues, a method of inhibiting madcham-mediated leukocyte to composition in a subject. The antibody, composition and methods are inflammatory boweld disease, Crohn's disease, primary bilary colitis, liver disease, primary bilary colitis, insulin dependent diabetes and coloral such account of the such antibody of the invention.
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                        Monoclonal antibody; mucosal addressin cell adhesion molecule; MAdCAM; inflammation; inflammatory bowel disease; Crohns disease; ulcerative colitis; diverticular disease; gastritis; liver disease; primary bilary cirrhosis; primary sclerosing cholangitis; insulin dependent diabetes; graft versus host disease; antiinflammatory; gastrointestinal-gen.; antiulcer; hepatotropic; antidiabetic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New antibody to Mucosal Adressin Cell Adhesion Molecule, useful for diagnosing and treating an inflammatory disease, e.g. inflammatory bo disease, ulcerative colitis, gastritis, insulin-dependent diabetes or
                                                                                                                                                                           Human monoclonal anti-MAdCAM antibody #9.
                                                                                                                                                                                                                     06-OCT-2005
                                                                                                                                                                                                                                                                                                       AEB45853 standard; protein; 469
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
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        antibody.
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Pred. No.
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                                                                                                                                                                                       CC nucleic acid molecule compositing a nuclebtide sequence encoding the heavy chain or its antigen-binding portion or the light chain or its antigen-compositing a nuclebtide sequence encoding the heavy composition of an antibody described above, a vector comprising the nucleic acid molecule, where the vector optionally comprises an composition control sequence operably linked to the nucleic acid molecule, a host cell comprising the vector optionally comprises and composition that specifically binds MACAM, a method of isolating an antibody or its antigen-binding corrien that specifically binds to MACAM, a method of isolating an antibody or its antigen-binding portion that specifically binds to MACAM, a composition portion that specifically binds to MACAM, a composition portion that specifically binds to MACAM, a composition and inhibits binding to expressing human MACAM, a method of inhibiting MACAM-mediated leukocyte composition that specifically binds to MACAM-mediated leukocyte composition that specifically binds to MACAM-mediated leukocyte composition into tissues, a method of inhibiting MACAM-mediated leukocyte composition and inhibits binding to cells composition and methods of inhibiting alphasheta7/MACAM-dependent cellular adhesion, cinflammation in a subject. The antibody, composition and methods are cuseful for disgnosing and treating inflammatory disease, e.g. cinflammatory bowel disease, Crohn's disease, ulcerative colitis, cinflammatory bowel disease, and the sequence represents a human monoclonal confirmation antibody of the invention.
                                                                            Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cell line that produces the human monoclonal antibody, a pharmaceutical composition comprising an amount of the monoclonal antibody or its antigen-binding portion and a pharmaceutical carrier, a method of treating inflammatory disease in a subject, an isolated cell line that produces the monoclonal antibody or its antigen-binding portion or the heavy chain or light chain of the antibody or of its portion, an isolated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to a human monoclohal antibody binding portion that specifically binds to mucosal acadesion molecule (MACCAM). The invention also relate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New antibody to Mucosal Adressin Cell Adhesion Molecule, useful for diagnosing and treating an inflammatory disease, e.g. inflammatory boundisease, ulcerative colitis, gastritis, insulin-dependent diabetes or
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                                                                                                                                                        Sequence
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TYYYYGMDV 131
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                                                                                                 Score 47;
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ATYYYFGLDV 11
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Pred. No. 0.5;
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The invention describes an isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a specific Clone where (I) binds to and neutralises botulinum neurotoxin type A (BONT/A). An isolated antibody (I) that specifically binds to an epitope Specifically bound by an antibody expressed by a clone chosen from clone CS 25, C35, 1C6, 3D12, B4, 1F3, huC25, Ar1, Ar2, WR1(V), WR1(T), 3-1, CC 3-8, 3-10 and ING1, where (I) binds to and neutralizes botulinum neurotoxin type A (BONT/A). Also described are: a polypeptide (II) CC comprising BONT/A neutralising epitope having an epitope that is specifically bound by an antibody expressed by clones as mentioned in (I); producing (I); and a composition (III) comprising several anticody in specific for a different epitope of a botulinum neurotoxin and the combination of comprising (I); and a kit comprising (I). (I) is useful for neutralising BONT/A antibody and for neutralisation than the single antibodies in surplus. (I) in surplus, where each of (I) is specific for a different epitope of the botulinum neurotoxin which combination of antibodies shows greater toxin neutralisation than the single antibodies in surplus. (I) is useful for diagnosing the botulism composition of involves contacting neurotoxin with (I) in surplus, where each of (I) is specific for a different epitope of the botulinum neurotoxin which composition of antibodies shows greater toxin neutralisation than the single antibodies in surplus. (I) is useful for diagnosing the botulism correction than the combination of antibodies associated with botulinum neurotoxin than the correction of diagnosis of botulism. This is the amino acid sequence of mouse heavy chain variable region complementarity determining region 3 (CDR3) from anti-botulinum neurotoxin antibodies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         antibacterial; antibody; botulinum neurotoxin type A; BONT/A; BONT/A neutralising epitope; anti-botulinum neurotoxin antibody; toxin neutralisation; botulinum neurotoxin poisoning; mouse;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New isolated antibody that neutralizes botulinum neurotoxin type A, useful for diagnosing botulism or for treating pathologies associated with botulinum neurotoxin poisoning.
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01-AUG-2002; 2002US-0400721P.
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                                                                                                                                   Sequence 13 AA,
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Marks JD,

heavy

ADR38712,

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The present invention relates to novel proteins (I) that bind to pregnancy-associated plasma protein A (PAPP-A ADW04676). (I) comprises a first and second immunoglobulin variable domain sequence which binds to PAPP-A. Also claimed are proteins (II) which comprise light chain (LC) and heavy chain immunoglobulin variable domain sequences which binds to PAPP-A. The proteins are useful for diagnosing, preventing or treating proliferative diseases such as glioblastoma, osteosarcoma and overgrowth of vascular smooth muscle cells following e.g., balloon angioplasty (which may cause restenosis). The proteins are especially useful for useful for treating diseases involving IGF regulated growth. The present sequence is one such immunoglobulin variable domain sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New pregnancy-associated plasma protein. (PAPP-A) binding proteins comprising immunoglobulin variable domain sequences, useful for diagnosing, preventing or treating diseases such as cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                proliferative disorder; restenosis; glioblastoma; osteosarcoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Cytostatic; Vasotropic; heavy chain variable domain;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PAPP-A immunoglobulin variable domain AB
Antibody; antibody engineering; antibody therapy; light chain variable region; heavy chain variable region;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example; SEQ ID NO 141; 168pp; English
                                                                                                                                                                                                                                                                                                                                                               ADZ41994 standard; peptide; 126 AA
                                                                                                                       variable region, B-CLL set II peptide #4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 46; DB 9;
Pred. No. 0.83;
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CC and heavy chain antibody genes, useful for treating B cell chronic CC lymphocytic leukemia (B-CLL). B-CLL is a disease of slowly proliferating CC CD5+ B lymphocytes. These cells express low levels of surface membrane Ig CC that serves as the receptor for antigen (BCR). Analysis of V region gene CC cassettes used by B-CLL clones differs from that in normal cells, with an CC increased frequency of VH3-07, VH4-34, and VH1-69 genes. This implies that the structure of the antibody molecule, and antigen specificity. CC play a role in the leukemic transformation of particular B cells. The CC patients with aggressive disease share the same classes of VH, D, JH, VL and JL antibody genes, forming sets of patients have a disorder selected from CH dogkin's disease, non-Hodgkin's lymphoma, Burkitt's lymphoma, myeloma or Systemic lupus erythematosus, myasthenia gravis, Grave's disease, type I CC diabetes mellitus, autoimmune peripheral neuropathy, and autoimmune Systemic lupus erythematosus, myasthenia gravis, Grave's disease, type I CC diabetes mellitus, autoimmune peripheral neuropathy, and autoimmune CC 13/JH5/VIkappa012/2/JIkappa1/kappa2 (Set II); VH3-CC 13/JH5/VIkappa012/2/JIkappa1/kappa2 (Set II); VH3-GC 13/JH6/VIlambda1c/JIlambda1 (Set V); VH1-03/D6-CC 13/JH5/VIkappa012/2/JIkappa1/kappa2 (Set VI); VH1-03/D6-CC 19/JH4/VIkappa012/2/JIkappa1 (Set VI); VH1-03/D6-CC 19/JH4/VIkappa012/2/JIkappa3 (Set VI); VH1-03/D6-19/JH4 (Set VId); VH5-05/D3-CC 19/JH4/VIkappa012/2/JIkappa3 (Set VII) Treating a patient having B-CLL CWith the above genes comprises administering an agent that binds to the antibody encoded by the antibody genes. The CC agent is an anti-idiotype antibody, a peptide antigen, or an aptamer. The CC apent sequence is an Ig H chain variable region, B-CLL set II peptide.
                                                                            Query Match
Best Local S
Matches 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Chronic lymphocytic leukemia; cytostatic; Hodgkins disease; lymphoma; Burkitts lymphoma; multiple myeloma; systemic lupus erythematosus; antiinflammatory; dermatological; immunosuppressive; myasthenia gravimuscular-gen.; neuroprotective; Graves disease; antithyroid; insulin dependent diabetes; diabetes mellitus; antidiabetic;
                                                                                                                                                                Sequence 126 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New isolated and purified preparation of light chain and heavy chain antibody genes, useful for diagnosing, preventing or treating B cell chronic lymphocytic leukemia, or in screening for agents that may treat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Fig
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Messmer BT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    autoimmune hemolytic anemia; antianemic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                new invention relates to combinations of light chain antibody genes
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105
                                    <u>بــا</u>
                                                                            Similarity
8; Conserv
                                    LATYYYFGLDV 11
LKRYYYYGLDV 115
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                                                                              Conservative
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                                                                              Score 46; DB Pred. No. 5.6; 1; Mismatches
                                                                                                  5.6;
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                                                                                                                     Length 126
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RESULT 14
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ADW04952
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                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention relates to novel proteins (1) that bind to pregnancy-associated plasma protein A (PAPP ADM04676). (1) comprises a first and second immunoglobulin variable domain sequence which binds to PAPP-A. Also claimed are proteins (II) which comprise light chain (LC) and heavy chain immunoglobulin variable domain sequences which binds to PAPP-A. The proteins are useful for disgnosing, preventing or treating proliferative diseases such as glioblastoma, osteosarcoma and overgrowth of vascular smooth muscle cells following e.g., balloon angioplasty (which may cause restenosis). The proteins are especially useful for useful for treating diseases involving IGP regulated growth. The present sequence is one such immunoglobulin variable domain sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New pregnancy-associated plasma protein-A (PAPP-A) binding proteins comprising immunoglobulin variable domain sequences, useful for diagnosing, preventing or treating diseases such as cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cytostatic; Vasotropic; proliferative disorder;
              Cytostatic; Vasotropic; heavy chain variable domain; proliferative disorder; restenosis; glioblastoma; os
                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 140 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unidentified
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADW04810 standard;
                                                                   PAPP-A immunoglobulin variable domain AB f03 heavy chain CDR3 SEQ ID 277.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example; SEQ ID NO 135; 168pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2005-080519/09.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Nixon A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        19-FEB-2003; 2003US-0448515P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         19-FEB-2004; 2004US-00783311.
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                                                                                                                                                                            ADW04952 standard; peptide;
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                                                                                                                                                                                                                                                                        109
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                                                                                                                                                                                                                                                                                                       LATYYYFGLDV 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hogan S;
                                                                                                                                                                                                                                                                                                                                           Conservative
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                                                                                                                                                                                                                                                                                                                                                            76.7%;
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                                                                                                                                                                                                                                                                                                                                                          Score 46;
Pred. No.
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                                                                                                                                                                                                                                                                                                                                               Mismatches
              glioblastoma; osteosarcoma
                                                                                                                                                                                                                                                                                                                                                              6.3;
                                                                                                                                                                                                                                                                                                                                                                               bB 9;
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Search completed: November 21, 2005, 12:20:00 Job time : 16.7271 secs
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                                                                                                                                             Query Match 75.0%; Score 45; DB 9; Length 16; Best Local Similarity 63.6%; Pred. No. 0.92; Matches 7; Conservative 3; Mismatches 1; Indels
                                                                                                                                                                                                                                                  The present invention relates to novel proteins (1) that bind to pregnancy-associated plasma protein A (PAPP-A ADW04676). (1) comprises a first and second immunoglobulin variable domain sequence which binds to PAPP-A. Also claimed are proteins (II) which comprise light chain (LC) and heavy chain immunoglobulin variable domain sequences which binds to PAPP-A. The proteins are useful for diagnosing, preventing or treating proliferative diseases such as glioblastoma, osteosarcoma and overgrowth of vascular smooth muscle cells following e.g., balloon angloplasty (which may cause restenosis). The proteins are especially useful for useful for treating diseases involving IGF regulated growth. The present sequence is one such immunoglobulin variable domain sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New pregnancy-associated plasma protein-A (PAPP-A) binding proteins comprising immunoglobulin variable domain sequences, useful for diagnosing, preventing or treating diseases such as cancer.
                                                                                                                                                                                                                      Sequence 16 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example; SEQ ID NO 277; 168pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2005-080519/09.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nixon A, Hogan S;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unidentified
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Listing first 45 summaries
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Perfect score:
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 Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.
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 protein search, using sw model
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Match
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60
1 LATYYYFGLDV 11
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1: pir1:*
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GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
                                                                                                                                                                                                                                                                       Length
 118
119
120
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128
132
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136
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627
627
581
126
375
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 PH1307
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PH1359
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spore germination
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Qy Db	RESULT 2 PH1361 Ig heavy chai C;Species: HC C;Date: 30-Se C;Accession: R;Wasserman, J. Exp. Med. A;Title: Pred A;Reference: A;Rocession: A;Molecule ty A;Residues: 1 A;Cressrefec C;Superfamily C;Keywords: h Query Match Best Local	RESULT 1 pH1307 Ig heavy ch C;Species: 1 C;Date: 30-: C;Accession R;Wasserman J. Exp. Med A;Title: Pr A;Reference A;Accession A;Molecule A;Accession A;Molecule C;Superfami C;Keywords: Query Mat Best Loca Matches Qy Db		444444 3321 4420 54321
LATYYYFGLDV 11      : :     LEPYYYYGMDV 18	m DJ region (clone C178-1 mo sapiens (man) p-193 #sequence_revision pH1361 176, 1577-1581, 1992 lominance of fetal type DJ number: PH1302; MUID:93094 pH1361 pH1361 pe: DNA -23 <was> -23 <was> rences: UNIPARC: UPI0000176 r. immunoglobulin V region leterotetramer; immunoglob similarity 63.6%; Pree 7; Conservative 2; N</was></was>	ain DJ region (clone C96-119) Homo Bapiens (man) Sep-1993 #Bequence_revision 3 : PH1307 edominance of fetal type DJH number: PH1302; MUID:9309476 : PH1307 type: DNA 1-19 < WASS arences: UNIPARC: UPI000017693 ly: immunoglobulin V region; heterotetramer; immunoglobulin che Similarity 66.7%; Pred. 6; Conservative 3; Mis 3 TYYYFGLDV 11 :     :     6 SYYYYGMDV 14	ALIGNMENTS ;	38 63.3 287 2 T27056 38 63.3 473 2 H86240 38 63.3 979 2 A39792 38 63.3 979 2 PH1601 37 61.7 14 2 PH1601 37 61.7 276 2 S16892 37 61.7 447 2 T09809 37 61.7 447 2 T09809 37 61.7 527 2 C70130 37 61.7 527 2 C70130 37 61.7 527 2 C72130 37 61.7 527 2 A42138 37 61.7 685 2 T22223 36.5 60.8 120 2 E49590 36.5 60.8 316 2 C75205 36 60.0 145 2 H97707
	gment) t_change 0 hane, S.; g children g children homology homology gth 23;	S.; ldre	MENT'S	hypothetical prote hypothetical prote transcription acti Ig H chain V-D-J r Ig heavy chain V-I probable transposa NADH2 dehydrogenas 91/cerol-3-phospha conditioned medium conditioned medium sel-1 protein - Ca Ig heavy chain V r hypothetical prote Ig heavy chain DJ 308 ribosomal prot
	7-May-1999 Rovera, G. with B precursor lympho	07-May-1999 Rovera, G. n with B precursor lympho		othetical prote inscription acti H Chain V-D-V r heavy chain V-I bable transposa H12 dehydrogenas H2 dehydrogenas cerol-3-phospha iditioned medium iditioned medium if protein - Ca heavy chain V r pothetical prote heavy chain DV r ribosomal prot

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C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 07-May-1999
C;Accession: PH1328
R;Masserman, R.; Galili, N.; Ito, Y.; Reichard, B.A.; Shane, S.; Rovera, G.
J. Exp. Med. 176, 1577-1581, 1992
A;Title: Predominance of fetal type DJH joining in young children with B precursor lymph A;Reference number: PH1302; MUID:93094761; PMID:1460419
A;Recession: PH1328
A;Molecule type: DNA
A;Residues: 1-29 < MASS-A;Residues: 1-29 < MAS
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S05271
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C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Date: 30-Jun-1992 #sequence_revision 30-Jun-1992 #text_change 31-Dec-2004
C;Accession: 805271; 804602
R;Kishimoto, T.
                               A;Molecule type: mRNA, A;Restdues: 1-160 <KISI>A;Restdues: 1-160 <KISI>A;COGE-references: UNIPROT:Q96BB9; UNIPARC:UPI0000176B50; A;CTCOGE-references: UNIPROT:Q96BB9; UNIPARC:UPI0000176B50; A;Kishimoto, T.; Okajima, H.; Okumoto, T.; Taniguchi, M. Nucletc Acide Res. 17, 4385, 1989
A;Telle: Nucleotide sequences of the cDNAs encoding the V-IA;Reference number: S04601; MUID:89296497; PMID:2500644
                                                                                                                                                                                                                                                                                                                                                                                         A; Reference number: S05270
A; Accession: S05271
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A;Residues: 1-160 <KLES
A;Cross-references: UNIPROT:O51659; UNIPARC:UPI00000575B2; GB:AE001171;
A;Experimental source: strain B31
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A;Authors: Smith, H.O.; Venter, J.C.
A;Title: Genomic sequence of a Lyme disease spirochaete,
A;Title: Genomic sequence of a Lyme disease spirochaete,
A;Reference number: A70100; MUID:98065943; PMID:9403685
A;Accession: D70189
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Ig heavy chain DJ region (clone C113-148) - human (fragment)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                submitted to the EMBL Data Library, March 1989
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A,Title: The complete genome of the hyperthermophilic bacterium Aquifex aeolicus. A,Reference number: A70300; MUID:98196666; PMID:9537320 A,Recession: E70380 A,Recession: P70380 A,Retatus: preliminary; nucleic acid sequence not shown; translation not shown A,Molecule type: DNA A,Molecule type: DNA A,Residues: 1-408 <AQF A,Residues: 1-408 <AQF A,Cross-references: UNIPROT:067072; UNIPARC:UPI00000564AB; GB:AE000714; NID:g2983446; A,Experimental source: strain VF5 C;Genetics:
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A; Residues: 1-144 < KIS2>
A; Cross-references: UNIPARC: UPI0000176B51; EMBL; X14584
A; Cross-references: UNIPARC: Upi0000176B51; EMBL; X14584
C; Superfamily: immunoglobulin homology
C; Keywords: heterotetramer; immunoglobulin
F; 1-19/Domain: signal sequence #status predicted <SIG>F; 20-160/Product: Ig heavy chain (fragment) #status pref; 34-117/Domain: immunoglobulin homology < IMM*
                                                                                                                                                                                R;Wasserman, R.; Galili, N.; Ito, Y.; Reichard, B.A.; Shane, S.; Rovera, G. J. Exp. Med. 176, 1577-1581, 1992
A;Title: Predominance of fetal type DJH joining in young children with B precursor lymphc A;Reference number: PH1302; MUID:93094761; PMID:1460419
A;Accession: PH1304
A;Ressidues: 1-19 <WAS>
A;Ressidues: 1-19 <WAS>
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N;Alternate names: Na+/H+ antiporter
C;Species: Aquifex aeolicus
C;Date: 08-May-1998 #sequence_revision 08-May-1998 #text_change 09-Jul-2004
C;Accession: E70380
                                                                                                                       A;Cross-references: UNIPARC:UPI0000176934
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
                                                                                                                                                                                                                                                                                                                               C;Species: Homo sapiens (man)
C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993
C;Accession: PH1304
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C;Superfamily: Aquifex aeolicus Na+/H+-exchanging
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116 IVSYYFFGLDL 126
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YYYFGLDV 11
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R; Wasserman, R.; Galili, N.; Ito, Y.; Reichard, B.A.; Shane, S.; Rovera, G. J. Exp. Med. 176, 1577-1581, 1992
A;Title: Predominance of fetal type DJH joining in young children with B precursor lymph A;Reference number: PH1302; MUID:93094761; PMID:1460419
A;Accession: PH1364
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A;Residues: 1-23 <WASZ
A;Cross-references: UNIPARC:UPI0000176949
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Molecule type: DNA
A;Residues: 1-19 <WAGA
A;Cross-references: UNIPARC:UPI000017690D
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: immunoglobulin
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A;Accession: $43960
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C;Species: Homo sapiens (man)
C;Date: 20-Oct-1994 #sequence_revision 26-Jul-1996 #text_change 17-Mar-1999
C;Accession: S43950
R;Wagner, S.D.; Williams, G.T.; Larson, T.; Neuberger, M.S.; Kitamura, D.;
Nucleic Acids Res. 22, 1389-1393, 1994
                                                                                                                                                                                             Ig heavy chain CDR3 region (clone 2-118B) - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 30-May-1997
C;Accession: PT025B
R;Yamada, M.; Wasserman, R.; Reichard, B.A.; Shane, S.; Caton, A.J.; Rovera, G.
J. Exp. Med. 173, 395-407, 1991
A;Title: Preferential utilization of specific immunoglobulin heavy chain diversity and A;Reference number: PT0222; MUID:91108337; PMID:1899102
A;Accession: PT025B
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                                                  A; Experimental source: B lymphocyte C; Superfamily: immunoglobulin V region; immunoglobulin homology C; Keywords: heterotetramer; immunoglobulin
                                                                                                                                                   A; Molecule type: DNA
A; Residues: 1-24 < YAM>
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C;Species: Homo sapiens (man)
C;Date: 30-Sep.1993 #sequence_revision 30-Sep-1993 #text_change 07-May-1999
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Similarity 75.0%;
6; Conservation
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Pred. No. 0.35;
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  42;
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Length 24;
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R.Wasserman, R.; Galill, N.; Ito, Y.; Reichard, B.A.; Shane, S.; Rovera, G. J. Exp. Med. 176, 1577-1581, 1992
A.Title: Predominance of fetal type DJH joining in young children with B precursor lymphc A;Reference number: PH1302; MUID:93094761; PMID:1460419
A;Accession: PH1371
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PH1371

Ig heavy chain DJ region (clone C111-145) - human (fragment)

C;Species: Homo sapiens (man)

C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 07-May-1999

C;Accession: PH1371
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Ig heavy chain DJ region (clone C100-136) - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 07-May-1999
C;Accession: PH1355
C;Accession: PH1355
R;Mortari, F.; Newton, J.A.; Wang, J.Y.; Schroeder Jr., H.W.
Eur. J. Immunol. 22, 241-245, 1992
Eur. J. Immunol. 22, 241-245, 1992
A;Title: The human cord blood antibody repertoire. Frequent usage of the V(H)7 gene famil A;Reference number: S26786; MUID:92111632; PMID:1730251
A;Accession: S26793
A;Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Cross-references: UNIPARC:UPI0000176945
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
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A;Title: Predominance of fetal type DJH joining in young children with B precursor lympho A;Reference number: PH1302; MUID:93094761; PMID:1460419
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A;Residues: 1-27 <MAS>
A;Cross-references: UNIPARC:UPI000017694B
C;Superfamily: immunoglobulin V region; in
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                                                                                                                                  Ig heavy chain V region - human (;Species: Homo sapiens (man) (;Species: Homo sapiens (man) (C;Date: 13-Jan-1995 #sequence_revision 13-Jan-1995 #text_change 20-Jun-2000 (;Date: 13-Jan-1995 #sequence_revision 13-Jan-1995 #text_change 20-Jun-2000 (;Accession: S26793
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A; Residues: 1-27 < WAS>
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YYYYGMDV 22
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Pred. No.
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Pred. No. 0.5;
2; Mismatches
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PH0961

Ig heavy chain V region (G6+ T-L33) - human (fragment)

C;Species: Homo sapiens (man)

C;Date: 17-Apr-1993 #sequence_revision 17-Apr-1993 #text_change 16-Aug-1996

C;Accession: PH0961

R;Martil, T.; Duffy, S.F.; Carson, D.A.; Kipps, T.J.

J. Exp. Med. 175, 983-991, 1992

A;Title: Evidence for somatic selection of natural autoantibodies.

A;Reference number: PH0952; MUID:92202880; PMID:1552291

A;Status: nucleic acid sequence not shown

A;Molecule type: DNA

A;Molecule type: DNA

A;Molecule type: DNA
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C;Species: Homo sapiens (man)
C;Date: 24-Feb-1994 #sequence_revision 24-Feb-1994 #text_change 16-Aug-1996
C;Accession: PH1666
R;Hillson, J.L.; Karr, N.S.; Oppliger, I.R.; Mannik, M.; Sasso, E.H.
J. Exp. Med. 178, 331-336, 1993
A;Title: The structural basis of germline-encoded VH3 immunoglobulin binding to staphylc A;Reference number: PH1642; MUID:93301610; PMID:8315388
A;Accession: PH1666
A;Molecule type: mRNA
A;Residues: 1-18 <HIL>
A;Cross-references: UNIPARC:UPI0000176BE7
A;Cross-references: UNIPARC:UPI0000176BE7
A;Cross-references: B cell
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
C;Keywords: heterotetramer; immunoglobulin
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A;Cross-references: UNIPARC:UPI0000115FCA; EMBL:X61019; NID:g32787; PIDN:CAA43353.1; PIDC:Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
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C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
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A; Residues: 1-119 < MAR>
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Pred. No. 2.2;
2; Mismatches
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Search completed: November 21, 2005, 12:22:14
Job time : 3.67949 secs

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      Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.
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Q7W919_BRAJA
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Q80W19_BRAJA
Q6AQ71_DESPS
Q6CMY2_HUMAN
Q96X97_SULTO
Q4FWX4_LEIMA
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Q6F8M6_ACIAD
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Q8DUZ3_ERWCT
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Q8DUZ3_ERALY
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Q4gal0 leishmania
Q8gnt9 bradyrhizob
Q74n07 nanoarchaeu
Q6gmy2 homo sapien
Q6x97 sulfolobus
Q4fwx4 leishmania
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	Score 44; DB 2; Length Pred. No. 60; 1; Mismatches 0; Ind	genome shotgun (WGS) 5.1; -; Genomic_DNA. WW; 474C3A40C23E8C5F	Y T.H., Lu C., Sun B., Pertea M., Allen J., tzhugh H.A., Morzaria EMBL/GenBank/DDBJ dat	T., de Villiers B., C A., Berriman M., Wilso g Z., Shallom S.J., We g Z., Shallom S.J., We	hite O.R., ; rva, a bovi	, de Villiers E.I ain A., Berriman Xiong Z <sup>*</sup> , Shallon oaibi A., Wasawo S.V., Creasy T.H blyum T.V., Perte	Apicomplexa; Piroplasmida; T	PRT; 1348 AA. Created) Last sequence update) Last annotation update)	Q9XTT1_CAEEL ALIGNMENTS	OGZEX6 ORYSA Q6F181 HUMAN Q8F175 LACPL LSP2 DROME Q7VQH6 CAMBE Q7VQH6 CAMBF Q7VQH6 CAMBF Q7VQH6 ANOGA Q5TWX6 ANOGA Q5GX03 HUMAN Q5TWX6 ANOGA Q5GX03 LEGPL Q5KZG3 RHOPA Q5KZG3 RHOPA Q6KSG3 RHOPA
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QAUFY8;
13-8EP-2005 (TrEMBLrel. 31, L:
13-8EP-2005 (TrEMBLrel. 31, L:
13-8EP-2005 (TrEMBLrel. 31, L:
13-8EP-2005 (TrEMBLrel. 31, L:
Hypothetical protein.
ORFNames=TA19710;
                                                                                                                                                                                                                                                                                                                                                STRAIN=ATCC 35210 / B31;

MEDLINE-98065943; PubMed-9403685; DOI=10.1038/37551;

MRDLINE-98065943; PubMed-9403685; DOI=10.1038/37551;

MRDLINE-98065943; PubMed-9403685; DOI=10.1038/37551;

MRDLINE-980685943; PubMed-9403685; DOI=10.1038/37551;

MAITING R., Clayton R.A., Doddon R.J., Hickey B.K., Gwinn M.L., Dougherty B.A., Tomb J.-F., Fleischmann R.D.,

Richardson D.L., Peterson J.D., Kerlavage A.R., Quackenbush J.,

Salzberg S.L., Hanson M., van Vugt R., Palmer N., Adams M.D.,

Salzberg S.L., Hanson M., van Vugt R., Palmer N., Adams M.D.,

Gocayne J.D., Weidman J.F., Utterback T.R., Watthey L., McDonald L.A.,

Artiach P., Bowman C., Garland S.A., Pujii C., Cotton M.D., Horst K.,

"Genomic sequence of a Lyme disease spirochaete, Borrelia

"Genomic sequence of a Lyme disease spirochaete, Borrelia
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01-JUN-1998 (TrEMBLrel. 24, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
Conserved hypothetical integral membrane protein.
OrderedLocusNames=BB0717;
Borrella burgdorferi (Lyme disease spirochete).
Bacteria; Spirochaetes; Spirochaetales; Spirochaetaceae; Borrelia;
Borrella burgdorferi group.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "The chromosome 1 sequence of Theileria annulata.";
Submitted (APR-2005) to the EMBL/GenBank/DDBJ datable EMBL; CR940347; CAIT4001.1; -; Genomic_DNA.

Hypothetical protein.
SEQUENCE 2262 AA; 262068 MW; FD9E8915243EF512 CI
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Pain A., Renauld H., Murphy L., Harris D.A.,
Hall N., Barrell B.G.;
                                                                                                                                                            PIR; D70189; D70189.

TIGR; BB0717; -.

Complete proteome; Hypothetical protein.

SEQUENCE 160 AA; 19300 MW; 9DE28A497C0F5737 CRC64;
                                                                                                                                                                                                                                                                        Nature 390:580-586(1997).
EMBL; AE001171; AAC67057.1; -;
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Eukaryota; Alveolata;
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Pred. No.
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Pred. No. 11;
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067072 AQUAE PRELIMINARY;
067072;
067072;
01-AUG-1998 (Tremblrel: 07, C
01-AUG-1998 (Tremblrel: 07, L
01-JUN-2003 (Tremblrel: 24, L
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Bacteria; Spirochaetes
Borrelia burgdorferi ç
NCBI TaxID=29519;
                                                                                               GO; GO:0016021; C:integral to membrane; ÉBA.
GO; GO:0015299; F:solute:hydrogen antiporter
GO; GO:0006885; P:regulation of pH; IBA.
GO; GO:0006810; P:transport; IEA.
InterPro; IPR006153; Na H Dorter.
Pfam; PF00999; Na H Exchanger; 1.
Complete proteome; Transmembrane; Transport.
SEQUENCE 408 AA; 43582 MW; 7E05B4B387A6C5
                                                                                                                                                                                                                                                            00 PIR
                                                                                                                                                                                                                                                                                                                                                                                                               MEDLINE=98196666; PubMed=9537320; DOI=10.1038/32831; Deckert G., Warren P.V., Gassterland T., Young W.G., Lenox A.L., Deckert G., Werbeek R., Snead M.A., Keller M., Aufay M., Huber Feldman R.A., Short J.M., Olsen G.J., Swanson R.V.; "The complete genome of the hyperthermophilic bacterium Aquifex
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Aquifax aeolicus.
Bacteria; Aquificae; Aquificales; Aquificaceae; Aquifex.
NCBI_TaxID=63363;
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Complete proteome; Hypothetical protein.

SEQUENCE 160 AA; 19293 MW; B65B15C5197BFB89 CRC64;
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25-OCT-2004 (TrEMBLrel. 28, Created)
25-OCT-2004 (TrEMBLrel. 28, Last sequence update)
25-OCT-2004 (TrEMBLrel. 28, Last annotation update)
Conserved hypothetical integral membrane protein.
OrderedLocusNames=BG0739;
                                                                                                                                                                                                                                                                                                          Nature 392:353-358(1998).
EMBL; AE000714; AAC07034.1; -; Genomic_DNA.
PIR; E70380; E70380.
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Name=napA2; OrderedLocusNames=AQ_929;
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                                                                                                                                                                                                                                                                                                                                                                                            "The complete genome of aeolicus.";
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Similarity 6; Conserv
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3; Mismatches
Score 43; DB
Pred. No. 28;
4; Mismatches
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RX MEDIINE-22827954; pubMed=12910271; DOI=10.1038/ng1227;

RA Parkhill J., Sebaihia M., Preston A., Murphy L.D., Thomson N.R.,

RA Parkhill J., Sebaihia M., Preston A., Murphy L.D., Thomson N.R.,

RA Harris D.B., Holden M.T.G., Churcher C.M., Bentley S.D., Mungall K.L.,

RA Achtman M., Atkin R., Baker S., Basham D., Bason N., Cherevach I.,

RA Achtman M., Atkin R., Baker S., Basham D., Bason N., Cherevach I.,

RA Achtman M., Atkin R., Baker S., Basham D., Bason N., Cherevach I.,

RA Chillingworth T., Collins M., Cronin A., Davis P., Doggett J.,

RA Feltwell T., Goble A., Hamlin N., Hauser H., Holroyd S., Jagels K.,

RA Feltwell T., Goble A., Hamlin N., Hauser H., Holroyd S., Jagels K.,

RA Leather S., Moule S., Norberczak H., O'Neil S., Ormond D., Price C.,

RA Rabbinowitsch E., Rutter S., Sanders M., Saunders D., Seeger K.,

RA Rabbinowitsch E., Rutter S., Sanders M., Saunders D.J.;

Unwin L., Whitehead S., Barrell B.G., Maskell D.J.;

"Comparative analysis of the genome sequences of Bordetella pertussis,

RT "Comparative analysis of the genome sequences of Bordetella pertussis,

RR BMSL, BX640415; CAB41776.1; -; Genomic_DNA.

DR GO; GO:0030288; C:periplasmic space (sensu Gram-negative Bact. . .; IEA.

BOG, GO:00308810; P:transport; IEA.
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Best Local S
Matches 7
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YVY71 BORPE
PRELIMINARY; PRT; 364 AA.
O7VY71
O1-OCT-2003 (TrEMBLrel. 25, Created)
O1-OCT-2003 (TrEMBLrel. 25, Last sequence update)
O1-OCT-2003 (TrEMBLrel. 26, Last annotation update)
O1-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Putative periplasmic solute-binding protein.
Name=smom; OrderedLocusNames=BP1487;
Bordetella pertussis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      07W926 BORPA PRELIMINARY; PRT; 364 AA. 07W926; 01-0CT-2003 (TrEMBLrel. 25, Created) 01-0CT-2003 (TrEMBLrel. 25, Last sequence update) 01-MAR-2004 (TrEMBLrel. 26, Last annotation update) Putative periplasmic solute-binding protein.
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InterPro; IPR006311; Tat.
InterPro; IPR004602; TRAP_transptDctP.
InterPro; IPR014602; TRAP_transptDctP.
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STRAIN=12822 / ATCC BAA-587;
MBDLINE=22827954; PubMed=12910271; DOI=10.1038/ng1227;
Parkhill J., Sebaihia M., Preston A., Murphy L.D., Thomson N.R.,
                                                                                                                                                                                                                                                                                                                                                   Name-smoM; OrderedLocusNames=BPP1948;
Bordetella parapertussis.
Bacteria; Proteobacteria; Betaproteobacteria; Burkholderiales;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TIGRFAMS; TIGRO1409; TAT_signal_seq; 1.
                                                                                                                                                       NUCLEOTIDE SEQUENCE.
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RMEDLINES-2887954; PubMed=12910271; DOI=10.1038/ng1227;

RA Parkhill J., Sebaihia M., Preston A., Mirphy L.D., Thomson N.R.,

RA Parkhill J., Sebaihia M., Preston A., Mirphy L.D., Thomson N.R.,

RA Harris D.E., Holden M.T.G., Churcher C.M., Bentley S.D., Mungall K.L.,

RA Achtman M., Atkin R., Baker S., Churcher K.D., Harris B., Quail M.A.,

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RA Chillingworth T., Collins M., Cromin A., Davis P., Doggett J.,

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RA Feltwell T., Goble A., Hamlin N., Hauser H., Holroyd S., Jagels K.,

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RA Feltwell T., Goble A., Hamlin N., Hauser H., Holroyd S., Jagels K.,

RA RABbinnowitsch E., Rutter S., Sanders M., Saunders D., Seeger K.,

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RA Abbinnowitsch E., Rutter S., Sanders M., Saunders D., Seeger K.,

RA Abbinnowitsch E., Rutter S., Sanders M., Saunders D.J.;

"Comparative analysis of the genome sequences of Bordetella pertussis,

RT Morth Bordetella parapertussis and Bordetella bronchiseptica.";

RMEL, BX640443; CAB32632.1; -; Genomic DNA.

DR EMBL, BX640443; CAB32632.1; -; Genomic DNA.

DR GO; GO:0006810; P:transport; IEA.
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"Comparative analysis of the genome sequences of Bordetella pertussis, Bordetella parapertussis and Bordetella bronchiseptica.";
Bordetella parapertussis and Bordetella bronchiseptica.";
BABL; BK640428; CAB37248.1; -; Genomic_DNA.
GO, GO.0030288; C.periplasmic space (sensu Gram-negative Bact. . .; IEJ
GO; GO:0006810; P:transport; IEA.
InterPro; IPR004631; Tat.
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01-OCT-2003 (TrEMBLrel. 25, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last sequence update)
Putative periplasmic solute-binding protein.
Name=smoM; OrderedLocusNames=BB3136;
Bordetella bronchiseptica (Alcaligenes bronchisepticus)
Bacteria; Proteobacteria; Betaproteobacteria; Burkholder
Alcaligenaceae; Bordetella.
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InterPro; IPR004682; TRAP_transptDctP.
Pfam; PF03480; SBP_bac_7; 1.
TIGRFAMs; TIGR01409; TAT_signal_seq; 1.
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            Score 42;
Pred. No.
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                                                                                                                                                                                                         8AB6672BA3BF09AD CRC64;
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                   38;
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Best Local S
Matches 7
                                                                          Bradyrhizobium japonicum USDA110.";
DNA Res. 9:189-197 (2002).
EMBL; BA000040; BAC49010.1; -; Genomic DNI
GO; GO:0030288; C:periplasmic space (sens)
GO; GO:005810; P:transport; IEA.
InterPro; IPR006311; Tat.
InterPro; IPR00631; TRAP_transptDctP.
Pfam; PF03480; SBP_bac_7; 1.
TIGRFAM8; TIGR01409; TAT_signal_seq; 1.
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QAQA10;
13-SEP-2005 (TrEMBLrel. 31, L
13-SEP-2005 (TrEMBLrel. 31, L
13-SEP-2005 (TrEMBLrel. 31, L
13-SEP-2005 (TrEMBLrel. 31, L
Hypothetical protein.
ORFNames=LmjF25.0840;
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Peascock C.S., Murphy L., Ivens A.C., Berriman M., Blackwell J.
Peascock C.S., Murphy L., Ivens A.C., Berriman M., Blackwell J.
Smith D., Collins M., Fosker N., Harris D., Oliver K., O'Neil
Saunders D., Seeger K., Warren T., Rajandream M., and Barrell
Saunders D., Seeger K., Warren T., Rajandream M., and Barrell
Submitted (JUN-2005) to the EMBL/GenBank/DDBJ databases.

BMBL, CT005264; CAJ04989.1; -; Genomic_DNA.
                                                                                                                                                                                                                                                                                                                                                                                                         MEDLINK=22464998, PubMed=12597275;
MEDLINK=22464998, PubMed=12597275;
Kaneko T., Nakamura Y., Sato S., Minamisawa K., Uchiumi T.,
Sasamoto S., Watanabe A., Idesawa K., Iriguchi M., Kawashima K.,
Sohara M., Matsumoto M., Shimpo S., Tsuruoka H., Wada T., Yamada
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    01-JUN-2003 (TrEMBLrel. 24, Created)
01-JUN-2003 (TrEMBLrel. 24, Last sequence update)
01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
Periplasmic mannitol-binding protein.
OrderedLocusNames=blr3745;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             QB9NT9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bradyrhizobium japonicum.
Bacteria, Proteobacteria, Alphaproteobacteria, Rhizobiales,
Bradyrhizobiaceae, Bradyrhizobium.
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Eukaryota; Euglenozoa;
EURI_TaxID=5664;
                                                                                                                                                                                                                                                                                                                                                             Tabata S.;
"Complete genomic sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRAIN=USDA 110;
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Pred. No.
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                                  23C264980A9F9B8D CRC64;
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RESULT 11
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Q74N07;
MEDLINE=22388257, PubMed=12477932; DOI=10.1073/pnas.242603899; MEDLINE=22388257, PubMed=12477932; DOI=10.1073/pnas.242603899; Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G., Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D., Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K., Hopkins R.F., Jordan H., Moore T., Max S.I., Mang J., Hsieh F., Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L., Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E., Stapleton M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C., Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J., Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H., Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
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Archaea; Nanoarchaeota; Nanoarchaeum
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InterPro; IPR003599; Ig-11ke.

InterPro; IPR003597; Ig-21.

InterPro; IPR003597; Ig_C1.

InterPro; IPR003596; Ig_MHC.

InterPro; IPR003596; Ig_W-C.

InterPro; IPR003596; Ig_W-C.

InterPro; IPR003596; Ig_W-C.

INTERPRO; IPR003596; Ig_W-C.

RAMART; SM00409; IG; 2.

R SMART; SM00406; IGC1; 4.

R SMART; SM00406; IGV; 1.

R SMART; SM00406; IGW; 1.

R PROSITE; PS00299; IG_MHC; UNKNOWN 3.

SEQUENCE 606 AA; 66185 MW; B6B38B51114E4C55 CRC64;
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                                                                                                                                                                                                                                                                                                          NUCLEOTIDE SEQUENCE.
STRAIN-JCM 10545 / 7;
MEDLINE=21456156; PubMed=11572479;
MEDLINE=21456156; PubMed=11572479;
Kawarabayasi Y., Hino Y., Horikawa H., Jin-no K., Takahashi M.,
Sekine M., Baba S.-I., Ankai A., Kosugi H., Hosoyama A., Fukui S.,
Nagai Y., Nishijina K., Otsuka R., Nakazawa H., Takamiya M., Kato
Yoshizawa T., Tanaka T., Kudoh Y., Yamazaki J., Kushida N., Oguchi
Aoki K.-I., Masuda S., Yanagii M., Nishimura M., Yamagishi A.,
Oshima T., Kikuchi H.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OrderedLocusNames=ST2615;
Sulfolobus tokodaii.
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01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-CCT-2003 (TrEMBLrel. 25, Last annotation update)
1062aa long hypothetical thermopsin.
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                                                                                                                                                                                              "Complete genome sequence of an aerobic thermoacidophilic Crenarchaeon, Sulfolobus tokodaii strain7.";
DNA Res. 8:123-140(2001).
EMBL; BA000023; BAB67731.1; -; Genomic_DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NCBI_TaxID=111955;
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                                                                                                                                      InterPro; IPR007981; Peptidase_A5. Pfam; PF05317; Thermopsin; 1.
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YYYYGMDV 142
                                                                                   proteome; Hypothetical protein.
1062 AA; 118024 MW; 166F4EE
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  70.0%;
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RX PubMed=16020728; DOI=10.1126/science.1112680;
RA Ivens A.C., Peacock C.S., Worthey E.A., Murphy L., Aggarwal G.,
RA Ivens A.C., Peacock C.S., Worthey E.A., Murphy L., Aggarwal G.,
RA Evriman M., Sisk E., Rajandream M.A., Adlem E., Aert R., Anupama A.,
RA Berriman M., Sisk E., Rajandream M.A., Adlem E., Aert R., Anupama A.,
RA Apostolou Z., Attipoe P., Bason N., Bauser C., Beck A., Beverley S.M.,
RA Apostolou Z., Attipoe P., Bason N., Bauser C.V., Collins M.,
RA Cadag E., Ciarloni L., Clayton C., Coulson R.M., Cronin A., Cruz A.K.,
RA Cadag E., Ciarloni L., Clayton C., Coulson R.M., Cronin A., Cruz A.K.,
RA Pavies R.M., De Gaudenzi J., Dobson D.E., Duesterhoeft A.,
RA Pazelina G., Fosker N., Frasch A.C., Fraser A., Fuchs M., Gabel C.,
RA Fazelina G., Fosker N., Harris D., Hertz-Powler C., Hilbert H., Horn D.,
RA Huang Y., Klages S., Knights A., Kube M., Larke N., Litvin L.,
RA Lord A., Louie T., Marra M., Masuy D., Matthews K., Michaeli S.,
RA Lord A., Louie T., Murler B., Rethardt R., Rieger M., Rinta J.,
RA Oliver K., O'Neil S., Pentony M., Pohl T.M., Price C., Purnelle B.,
RA Oliver K., O'Neil S., Pentony M., Pohl T.M., Price C., Purnelle B.,
RA Oliver M., Rabbinowitsch E., Reinhardt R., Rieger M., Rinta J.,
RA Robben J., Robertson L., Ruiz J.C., Rutter S., Saunders D.,
RA Schafer M., Schein J., Schwartz D.C., Séeger K., Seyler A., Sharp S.,
RA Volckaert G., Wambut R., Warren T., Woodward J., Zhou S.,
RTM Common C. the Kinerorlastid Darratte Letenania mater ""
"The Crowne C. the Kinerorlastid Darratte Letenania mater"
"The Crowne C. the Kinerorlastid Darratte Letenania mater"
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QAFWX4;
13-SEP-2005 (TrEMBLrel. 31,
13-SEP-2005 (TrEMBLrel. 31,
13-SEP-2005 (TrEMBLrel. 31,
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EMBL; CP000081; AAZI4369.1;
Hypothetical protein.
SEQUENCE 3212 AA; 342958
                                                                                                                     Desulfotalea psychrophila.
Bacteria; Proteobacteria; Deltaproteobacteria; Desulfobacterales;
Desulfobulbaceae; Desulfotalea.
NCBI_TaxID=84980;
                                                                                                                                                                                                                                                                                                                                                                                       QGAQZI DESPS PRELIMINARY; PRT; 716 AA. QGAQZI, QGAQZI, 28, Created) 25-OCT-2004 (TrEMBLrel. 28, Last sequence update) 25-OCT-2004 (TrEMBLrel. 28, Last annotation update) 25-OCT-2004 (TrEMBLrel. 28, Last annotation update)
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Eukaryota; Euglenozoa;
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    NUCLEOTIDE SEQUENCE.
STRAIN=LSv54 / DSM 12343;
                                                                                                                                                                                                                                                                                                                                                            Related to mercuric reductase.
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RA PubMed=15305914; DOI=10.1111/j.1466-2920.2004.00665.x;
RA Rabuser M., Zibat A., Lombardot T., Becker I., Amann J., Gellner K.,
RA Teeling H., Leuschner W.D., Gloeckner F.-O., Lupas A.N., Amann R.,
RA Klenk H.-P.;
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RA Klenk H.-P.;
RA G.;
RO:GO:GO050601; Pedictase Activity; IEA.
RA G.;
RO:GO:GO050601; P.; PAD binding; IEA.
RA G.;
RA G.;
RA RA Teeling H.-P.
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GenCore version 5.1.6 Copyright (c) 1993 - 2005 Compugen Ltd.

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Result
No.
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Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Minimum DB seq length: 0
Maximum DB seq length: 200000000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Scoring table:
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Perfect score:
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      Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.
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      protein search, using sw model
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Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 November 21, 2005, 11:49:31; Search time 4.15018 Seconds (without alignments) 219.131 Million cell updates/sec
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1: /cgn2_6/ptodata/1/1aa/5_COMB.pep:*

2: /cgn2_6/ptodata/1/1aa/6_COMB.pep:*

3: /cgn2_6/ptodata/1/1aa/H_COMB.pep:*

4: /cgn2_6/ptodata/1/1aa/RE_COMB.pep:*

5: /cgn2_6/ptodata/1/1aa/RE_COMB.pep:*

6: /cgn2_6/ptodata/1/1aa/RE_COMB.pep:*
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US-08-053-131-78
US-08-045-641-78
US-08-08-645-78
US-08-096-762-78
US-08-09-308-865-78
US-09-042-353-78
US-09-047-625A-84
US-09-047-625A-84
US-09-08-3131-73
US-08-053-131-73
US-08-053-131-73
US-08-053-1408B-73
US-08-053-408B-73
US-08-053-131-83
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                                                                                                                                                                                                                                                       Sequence 78, Ay Sequence 78, Ay Sequence 78, Ay Sequence 275, Ay Sequence 78, Ay Sequence 78, Ay Sequence 78, Ay Sequence 84, Ay Sequence 73, Ay Sequence 74, 
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78, Appl
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78, Appl
78, Appl
275, Appl
84, Appl
84, Appl
84, Appl
73, Appl
83, Appl
84, Appl
84, Appl
84, Appl
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US-08-053
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45	44	43	42	41	40	9	38	37	36	35	34	ည	32	31	30	29	28	
42	42	42	42	42	42	42	42	42	42	42	42	42	42	42	42	42	42	
70.0	70.0	70.0	70.0	70.0	70.0	70.0	70.0	70.0	70.0	70.0	70.0	70.0	70.0	70.0	70.0	70.0	70.0	
236	236	236	236	167	128	128	128	128	119	119	36	36	36	36	36	36	36	
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US-09-453-234-104	US-09-453-234-64	US-09-456-090A-104	US-09-456-090A-64	US-09-472-087-80	US-09-497-625A-79	US-09-497-625A-77	US-09-840-459-79	US-09-840 #459-77	US-09-497-625A-88	US-09-840-459-88	PCT-US92-10983-84	US-08-758-417A-129	US-09-042-393-281	US-08-308-805-84	US-08-096-762-84	US-07-853-408B-84	US-08-645-641-84	
	Sequence 64, Appl	104	64,	Sequence 80, Appl	-	-		-	9 6	, a	4	129,	787	4	24	4.		

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TELEFAX: 415-326-2422
; INFORMATION FOR SEQ ID NO: 7
; SEQUIENCE CHARACTERISTICS:
; LENGTH: 33 amino acids
TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
US-08-645-641-78
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Best Local Similarity
%** 6; Conserva
                                                                                                                                 US-07-853-408B-78
                                                                                      sequence 78, Application US/07853408B
Patent No. 5789650
                                                                                                                                                                                                                                                                                   Query Match 71.7%;
Best Local Similarity 66.7%;
Matches 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/08/645,641
FILING DATE: 20 MAY-1996
CLASSIFICATION: 800
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/904,068
FILING DATE: 23 JUN-1992
ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M.
REGISTRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 14643-000913
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   -08-645-641-78
                                                                          GENERAL INFORMATION:
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICANT: Lonberg, Nils
APPLICANT: Kay, Robert M.
TITLE OF INVENTION: Trans
TITLE OF INVENTION: Produ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Lonberg, Nils
APPLICANT: Kay, Robert M.
TITLE OF INVENTION: Transgenic
TITLE OF INVENTION: Producing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
ADDRESSEE: William M. Smith
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ZIP: 94111-3834
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                                                                                                                                                                                                          10 SYYYYGMDV 18
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66.7%;
Transgenic No. 5789650-Human Animals for Producing Heterologous Antibodies
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 Mismatches

                                                                                                                                                                                                                                                                                                       Score 43; DB 1; Length 33; Pred. No. 1.6;
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Pred. No. 1.6;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-07-853-408B-78
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: Patentin Release #1.0, Version #1.25
SOPTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/096,762
FILING DATE: 22-UUL-1993
CLASSIFICATION: 800
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/053,131
FILING DATE: 26-APR-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/990,860
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 78, Application US/08096762
Patent No. 5814318
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TBLEPHONE: 415-326-2400
TBLEFAX: 415-326-2422
INFORMATION FOR SEQ ID NO: 7
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION UNMBER: US/07/853,408B
FILING DATE: 1990318
CLASSIFICATION: 800
CLASSIFICATION: 800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
NAME: SMILL, WILLIAM M.
REGISTRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 14643-9
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 210
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend Khourie and Crew STREET: One Market Plaza, Steuart Tower, Suite 200
                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Lonberg, Nils
APPLICANT: Kay, Robert M.
TITLE OF INVENTION: Transgenic No. 5814818-Human Animals for
TITLE OF INVENTION: Producing Heterologous Antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 33 amino acids
TYPE: AMINO ACID
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STREET: One Market I
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Local Similarity 66.7%;
les 6; Conservative
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                                                                                                                                                                                                                                                                                                         ZIP: 94105
                                                                                                                                                                                                                                                                                                                                 COUNTRY:
                                                                                                                                                                                                                                                                                                                                                   CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE:
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One Market Plaza, Steuart Tower, Suite 2000
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Pred. No. 1.6;
3; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 1; Length 33;
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US-08-096-762-78
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SEQUENCE CHARACTERISTICS:
LENGTH: 33 amin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Kay, Nils.
APPLICANT: Kay, Robert M.
TITLE OF INVENTION: Transgenic No. 5877397-
TITLE OF INVENTION: Producing Heterologous
NUMBER OF SEQUENCES: 150
        PILING DATE: 23-JUN-1992
ATTORNEY/AGENT INFORMATION:
NAME: SMAIL, William M.
REGISTRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 1464
TELECOMUNICATION INFORMATION:
TELEPHONE: 415-326-2400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       quence 78, Application US/08308865
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
NAME: SMith, William M.
REGISTRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 14643-9-4
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-326-2400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: linear MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: 18-MAR-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0:
FILING DATE: 17-DEC-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                           MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC COMPANIA
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/308,865
                                                                                                                                                                            FILING DATE: 800 CLASSIFICATION: 800 PRIOR APPLICATION DATA: APPLICATION NUMBER: (
                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
ADDRESSEE: William M. Smith
STREET: One Market Plaza, Steuart Tower, Suite 2000
CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 33 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 16-DEC-1992
IOR APPLICATION DATA:
APPLICATION NUMBER: US 0:
FILING DATE: 23-JUN-1992
                                                                                                                         FILING DATE: APPLICATION NUMBER: US 07/904,068 FILING DATE: 23-JUN-1992
                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 94105
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                                                                                                                                                                                                                                                                                                                                                                                                                         USA
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66.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Transgenic No. 5877397-Human Animals for Producing Heterologous Antibodies
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                                                14643-9-1-1
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Pred. No. 1.6;
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; MOLECULE TYPE: peptide
US-08-308-865-78
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Matches 6; Conserv
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SEQUENCE CHARACTERISTICS:
LENGTH: 33 amino acids
TYPE: amino acid
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APPLICATION NUMBER: US/0

FILING DATE: 13-MAR-1998

CLASSIFICATION: 800

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 0
                                                                                                                                                                                                                                                                                                                                                                                                                         PILING DATE: 17-DEC-19
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US
PILING DATE: 18-MAR-19
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US 07/904,068
FILING DATE: 23-JUN-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/990,860
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MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Lonberg, Nils
APPLICANT: Kay, Robert M.
TITLE OF INVENTION: Transgenic No. 6259458-Human Animals
TITLE OF INVENTION: Producing Heterologous Antibodies
                                                                                                                                                APPLICATION NUMBER: US 01
PILING DATE: 22-JUL-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 01
FILING DATE: 18 NOV-1993
                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
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                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 01
PILING DATE: 03-DEC-1993
                                                                                                                                                                                                                                                 FILING DATE: 26-APR-1993 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
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                                      10-DEC-1993
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                                                        US 08/165,699
                                                                                                               US 08/161,739
                                                                                                                                                                        US 08/155,301
US 08/209,741
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Pred. No. 1.6;
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:

US 08/352,322

FILING DATE: 09-MAR-1994

APPLICATION NUMBER: US 08/544,464 FILING DATE: 10-OCT-1995 APPLICATION NUMBER: US 08/352,322

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; TOPOLOGY: linear; MOLECULE TYPE: peptide US-09-042-353-275
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  S-08-758-417A-123
Sequence 123, Application US/08758417A
Patent No. 6300129
GENERAL INFORMATION:
APPLICANT: Lonberg, Nils
Kay, Robert M.
Kay, Robert M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity
Matches 6; Conserv
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TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 275
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: Apple, Randolph T.
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 014643-009040US
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/728,463
FILING DATE: 10-0CT-1996
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: WO PCT/US97/21803 FILING DATE: 01-DEC-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US 08/758,417 FILING DATE: 02-DEC-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: WO PCT/US96/16433 FILING DATE: 10-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 07-DEC-1994
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/758,417A
APPLICATION NUMBER: US/08/758,417A
FILING DATE: 02-Dec-1996
CLASSIFICATION: «Unknown»
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/728,463
FILING DATE: 10-OCT-1996
                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 417
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: Transgenic No. 6300129-Human Animals for Producing Heterologous Antibodies
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                                                                                                                                                                                                                                                                                                          CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                      COUNTRY: USA
                                                                                                                                                                                                                                                                    ZIP: 94111-3834
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66.7%;
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Pred. No. 1.6;
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PCT-US92-10983-78
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 78, Application PC/TUS9210983 GENERAL INFORMATION:
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                                                                                                                                                 COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 152
CORRESPONDENCE ADDRESS:
ADDRESSEE: William M. Smith
ADDRESSEE: One Market Plaza, Steuart Tower, Suite 2000
STREET: One Market Plaza, Steuart Tower, Suite 2000
    ATTORNEY/AGENT INFORMATION:
NAMS: SMICH, WILLIAM M.
REGISTRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Lonberg, Nils
APPLICANT: Kay, Robert M.
TITLE OF INVENTION: Transgenic Non-Humah An
TITLE OF INVENTION: Producing Heterologous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
                                                                                                             APPLICATION NUMBER: P
FILING DATE: 19921217
                                                                                           CLASSIFICATION:
                                                                                                                                                                                                                                                                                                            COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear
MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 123:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: 16-DEC-19
ATTORNEY/AGENT INFORMATION:
NAME: Serafini, Andrew
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            10 SYYYYGMDV 18
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                                                                                                                                                                                                                                                                                                                                                       San Francisco
                                                                                                                                                                                                                                                                                                                                  California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REGISTRATION NUMBER: 41,303
REFERENCE/DOCKET NUMBER: 014643-009030US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRANDEDNESS: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US 07/990,860 FILING DATE: 16-DEC-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE: 10-DEC-
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 22-JUL-1993
APPLICATION NUMBER: US 08/053,131
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US 08/165,699
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US 08/209,741
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TLING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ILING DATE: 26-APR-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PPLICATION NUMBER: US 08/096,762
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   'ILING DATE: 18-NOV-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ILING DATE: 03-DEC-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ILING DATE: 09-MAR-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 71.7%; Score 43; DB 2; Length 33; 66.7%; Pred. No. 1.6; tive 3; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Transgenic Non-Humah Animals for
                                                                                                                                      PCT/US92/10983
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Andrew T.
        14643-9-2
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Gaps

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US-09-840-459-84
                                                                                                                                                                                             RESULT 10
US-09-497-625A-84
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; ORGANISM: Homo sapiens
US-09-840-459-84
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/09/840,459
CURRENT FILING DATE: 2001-02-02
PRIOR APPLICATION NUMBER: PCT/US01/03537
PRIOR FILING DATE: 2001-02-02
PRIOR APPLICATION NUMBER: 09/497,625
PRIOR APPLICATION NUMBER: 09/497,625
PRIOR APPLICATION NUMBER: 09/359,193
PRIOR APPLICATION NUMBER: 09/359,193
PRIOR APPLICATION NUMBER: 09/3193
PRIOR FILING DATE: 1998-07-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 84, Application US/09840459 Patent No. 6696550
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                                                                                                                          Sequence 84, Application US/09497625A Patent No. 6727349
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                               Matches
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SEQ ID NO 84
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Horvath, Christopher
APPLICANT: Newman, Walter
APPLICANT: Jones, S. Tarran
APPLICANT: O'Brien, Siobhan H.
APPLICANT: O'Keefe, Theresa
APPLICANT: O'Keefe, Theresa
TITLE OF INVENTION: HUMANIZED ANTI-CCR2 ANTIBODIES AND
TITLE OF INVENTION: METHODS OF USE THEREFOR
FILE REFERENCE: 1855.1052-012
APPLICANT: LaRosa, Gregory J.
APPLICANT: Horvath, Christopher
APPLICANT: Newman, Walter
APPLICANT: Jones, S. Tarran
APPLICANT: O'Brien, Siobhan H.
APPLICANT: O'Keefe, Theresa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: LaRosa, Gregory J.
APPLICANT: Horvath, Christop
APPLICANT: Newman, Walter
APPLICANT: Jones, S. Tarran
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: peptide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: AMINO ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY:
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                                                                                                                                                                                                                                                                                       106 SYYYYGMDV 114
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                                                                                                                                                                                                                                                                                                                                   3 TYYYFGLDV 11
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nilarity 66.7%;
Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 43; DB 4;
Pred. No. 1.6;
                                                                                                                                                                                                                                                                                                                                                                                                  Score 43; DB 2;
Pred. No. 6.4;
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                                                                                                                                                                                                                                                                                                                                                                               0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION UMBER: US/08/053,131

PILING DATE: 26-APR-1993

CLASSIFICATION BOTA: 800

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 07/990,860

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 07/990,860

PILING DATE: 16-DEC-1992
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: HUMANIZED ANTI-CCR2 ANTIBODIES AND TITLE OF INVENTION: METHODS OF USE THEREFOR FILE REPERENCE: 1855.1052-004

CURRENT PAPALICATION NUMBER: US/09/497,625A

CURRENT FILING DATE: 2000-02-03

PRIOR APPLICATION NUMBER: 09/359,193

PRIOR PILING DATE: 1999-07-22

PRIOR APPLICATION NUMBER: 09/121,781

PRIOR PILING DATE: 1998-07-23

NUMBER: 09 SEQ ID NOS: 106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      sequence 73,
                                                            TELEFAX: 415-326-2422
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/810
FILING DATE: 17-DEC-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/853
FILING DATE: 18-MAR-1992
ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M.
REGISTRATION NUMBER: 30,223
REGISTRATION NUMBER: 30,223
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 12
TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend Khourie and Crew STREET: One Market Plaza, Steuart Tower, Suite 200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Lonberg, Nils
APPLICANT: Kay, Robert M.
TITLE OF INVENTION: Transgenic No. 5661016-Human Animals for
TITLE OF INVENTION: Producing Heterologous Antibodies
                                                                                                                                        TELECOMMUNICATION INFORMATION: TELEPHONE: 415-326-2400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Y Match 71.7%;
Local Similarity 66.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STATE: California
                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: 14643-9-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY:
                                                                                                                                      TELEPHONE:
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                         : 29 amino acids
amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     USA
                                                                                                                                                                                                                                                                                              US 07/853,408
                                                                                                                                                                                                                                                                                                                                                               US 07/810,279
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Sequence 73, Application US/07853408B; Patent No. 5789650; GENERAL INFORMATION:
                                                                                                                                                                                            Query Match
Best Local Similarity
""" hes 6; Conserve
                                                                    US-07-853-408B-73
                                                                                       RESULT 13
                                                                                                                                           밁
                                                                                                                                                                                                                                                                              , MOLECULE TYPE: peptide US-08-645-641-73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; MOLECULE TYPE: peptide US-08-053-131-73
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Best Local 9
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APPLICATION NUMBER: US 07/904,068

PILING DATE: 23-JUN-1992

ATTORNEY/AGENT INFORMATION:

NAME: Smith, William M.

REGISTRATION NUMBER: 30,223

REFERENCE/DOCKET NUMBER: 14643-0009
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
APPLICANT: Lonberg, Nils
APPLICANT: Kay, Robert M.
                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: 415-326-2422
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 29 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/645,641
FILING DATE: 20-MAY-1996
CLASSIFICATION: 800
 APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION: TELEPHONE: 415-326-2400
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
ADDRESSEE: William M. Smith
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: Transgenic No. 5719032-Human Animals TITLE OF INVENTION: Producing Heterologous Antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        y Match 70.0%;
Local Similarity 75.0%;
hes 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET: Two Embarcadero Center, 8th Floor CITY: San Francisco STATE: California
                                                                                                                                                                                                                                                                                                                STRANDEDNESS: single TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                             LENGTH: 29 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEPHONE:
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YYYYGMDV 14
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YYYYGMDV 14
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Lonberg, Nils
                                                                                                                                                                                                              Conservative
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                                                                                                                                                                                                                             Score 42; [
Pred. No. 2
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                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                           DB 1; Length 29;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M.
NAME: Smith, William M.
REGISTRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 1464.
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-326-2400
TELEPHONE: 415-326-2422
INFORMATION FOR SEQ ID NO: 73:
SEQUENCE CHARACTERISTICS:
LENGTH: 29 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Loc
Matches
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIAN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/096,762
FILING DATE: 22-UTL-1993
CLASSIFICATION: 800
PRICR APPLICATION DATA:
APPLICATION NUMBER: US 08/053,131
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Patent No. 5814318
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 73,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                      APPLICANT: Lonberg, Nils
APPLICANT: Kay, Robert M.
APPLICANT: Kay, Robert M.
TITLE OF INVENTION: Transgenic No. 5814318-Human Animals for
TITLE OF INVENTION: Producing Heterologous Antibodies
NUMBER OF SEQUENCES: 210
CORRESPONDENCE ADDRESS:
ADDRESSES: Townsend and Townsend Khourie and Crew
ADDRESSES: Townsend and Townsend Khourie and Crew
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/853,408B
FILLING DATE: 1920318
CLASSIFICATION: 800
CLASSIFICATION: 800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Kay Robert M.

TITLE OF INVENTION: Transgenic No. 5789650-Human Animals for TITLE OF INVENTION: Producing Heterologous Antibodies NUMBER OF SEQUENCES: 150

CORRESPONDENCE ADDRESS: CORRESPONDENCE ADDRESS: William M. Smith STREET: One Market Plaza, Steuart Tower, Suite 2000

CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                  STREET: One Market F
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity
nes 6; Conserv
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                                                                                                                                                                                                                                            ZIP: 94105
                                                                                                                                                                                                                                                              COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY:
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One Market Plaza, Steuart Tower, Suite 200
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75.0%;
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Pred. No. 2.1;
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FILING DATE: 26-APR-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0
FILING DATE: 16-DEC-1992

US 07/990,860

PRIOR APPLICATION DATA:

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RESULT 15
US-08-308-865-73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE: peptide US-08-096-762-73
  COUNTRY: USA
ZIP: 94105
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/308,865
FILING DATE:
APPLICATION NUMBER: US/08/145,707
FILING DATE:
APPLICATION NUMBER: US 07/904,068
FILING DATE: 23-JUN-1992
ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M.
REGISTRATION NUMBER: 30,223
ATTORNEY Smith, WILLIAM M.
REGISTRATION NUMBER: 1,223
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE: 415-326-2400
TELEPAX: 415-326-2422
INPORMATION FOR SEQ ID NO: .73:
SEQUENCE CHARACTERISTICS:
LENGTH: 29 amino acids
TYPE; amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 73, Application US/08308865
Patent No. 5877397
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 70.0%;
Best Local Similarity 75.0%;
Matches 6; Conservative
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APPLICATION NUMBER: US 07/810,279

PILING DATE: 17-DEC-1991

ATTORNEY/AGENT INFORMATION:

NAME: Smith, William M.

REGISTRATION NUMBER: 30,223

REFERENCE/DOCKET NUMBER: 14643-9-4

TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
APPLICANT: Lonberg, Nils
APPLICANT: Kay, Robert M.
TITLE OF INVENTION: Transgenic No. 5877397-Human Animals for
TITLE OF INVENTION: Producing Heterologous Antibodies
NUMBER OF SEQUENCES: 150
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/853,408
FILING DATE: 18-MAR-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: William M. Smith
STREET: One Market Plaza, Steuart Tower, Suite 2000
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US 07/904,068 FILING DATE: 23-JUN-1992
REFERENCE/DOCKET NUMBER: 14643-9-1-1
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7 YYYYGMDV 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Indels
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SEQUENCE CHARACTERISTICS:
LENGTH: 29 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-308-865-73
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                   ঠ
                                               Best Loc
Matches
                                                                         Query Match
                                                                                                                                                                                       INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-326-2400
                                           y Match 70.0%;
Local Similarity 75.0%;
les 6; Conservative
 YYYYGMDV 14
                      YYYFGLDV 11
                                                   Mismatches
                                                              Score 42; DB Pred. No. 2.1;
                                                                           DB 1,
                                                                          Length 29;
                                                    0; Indels
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                                                    Gaps
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Search completed: November 21, 2005, 12:07:38 Job time : 5.15018 secs

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GenCore version 5.1.6 Copyright (c) 1993 - 2005 Compugen Ltd.

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Regult
No.
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Maximum DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Run
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Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence:
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    Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
                                                                                                                                                                                                                                                                                                                                        Score
                                                                                                                                                                                                                                                                                                                                                                                                 is derived by analysis of the total score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            seq length: 0
seq length: 2000000000
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    99
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BLOSUM62
Gapop 10.0 , Gapext 0.5
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60
   November 21, 2005, 11:50:51;
                                                                                                                                                                                                                                                                                                                                        Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1867569 seqs, 417829326 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                              Published Applications AA Main:*

1: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*

2: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

3: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

4: /cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*

5: /cgn2_6/ptodata/1/pubpaa/US11A_PUBCOMB.pep:*

6: /cgn2_6/ptodata/1/pubpaa/US11A_PUBCOMB.pep:*
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    US-10-632-706-126
US-10-633-706-127
US-11-633-706-127
US-11-631-485-15
US-10-452-593-101
US-10-452-593-15
US-11-031-485-18
US-11-031-485-18
US-11-031-485-18
US-10-632-706-114
US-10-783-311-137
US-10-783-311-137
US-10-783-311-271
US-10-783-311-271
US-10-783-311-271
US-10-783-311-271
US-10-783-311-271
US-10-293-418-1600
US-10-293-418-1600
US-10-293-418-2876
US-10-293-418-2876
US-10-293-418-2876
US-10-293-418-2876
US-10-293-418-2876
US-10-293-418-2876
US-10-293-418-2876
US-10-293-418-2876
US-10-293-418-2876
US-10-293-418-2876
US-10-309-762-18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; Search time 13.619 Seconds (without alignments) 337.478 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                    distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1867569
Sequence 126, App Sequence 117, App Sequence 110, App Sequence 101, App Sequence 15, Appl Sequence 15, Appl Sequence 15, Appl Sequence 114, App Sequence 114, App Sequence 114, App Sequence 277, App Sequence 277, App Sequence 277, App Sequence 271, App Sequence 272, App Sequence 273, App Sequence 274, Appl Sequence 2876, App Sequence 2876, App Sequence 2876, App Sequence 2876, App Sequence 18, Appl Sequence 18, Appl Sequence 19, Appl
                                                                                                                                                                                                                                                                                                                                         Description
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RESULT 1  US-10-632-706-126 ; Sequence 126, Application US/10632706 ; Publication No. US20040175385A1 ; GENERAL INFORMATION: ; APPLICANT: MARKS, JAMES D. ; APPLICANT: MARKS, JAMES D. ; TITLE OF INVENTION: THERAPEUTIC MONOCLONA TITLE OF INVENTION: NEUROTOXINS ; FILE REFERENCE: 407T-895120US ; CURRENT APPLICATION NUMBER: US/10/632,706 ; CURRENT FILING DATE: 2003-08-01 ; PRIOR APPLICATION NUMBER: US 60/400,721 ; PRIOR APPLICATION NUMBER: US 09/144,806 ; PRIOR FILING DATE: 1998-08-31 ; NUMBER OF SEQ ID NOS: 278 ; SOFTWARE: PATENTIN 11 TYPE: PRT ORGANISM: Artificial		28 44 73 31 44 73 35 44 73 36 44 73 37 44 73 38
P-706-126  126, Application US/1063  Lion No. US20040175385A1  ANT: MARKS, JAMES D.  ANT: MARKS, JAMES D.  INFORMATION: THERAPEUTIC OF INVENTION: NEUROTOXINS REFERRICE: 407T-895120US UT APPLICATION UMBER: US 107  TILING DATE: 2003-08-01 APPLICATION NUMBER: US 60/ FILING DATE: 2002-08-01 APPLICATION NUMBER: US 09/ FILING DATE: 1998-08-31 APPLICATION NUMBER: US 09/ FILING DATE: 1998-08-31  APPLICATION NUMBER: US 09/ FILING DATE: 1998-08-31  APPLICATION NUMBER: US 09/ FILING DATE: 1998-08-31  APPLICATION NUMBER: US 09/ FILING DATE: 1998-08-31  FILING DATE: 1998-08-31  ROF SEQ ID NOS: 278  NO 126  HN 0126  HN 01		.3 1127 .3 1277 .3 127
ULT 1 10-632-706-126 equence 126, Application US/10632706 equence 126, Application US/10632706 ublication No. US20040175385A1 ENERAL INFORMATION: APPLICANT: MARKS, JAMES D. APPLICANT: MARKS, JAMES D. TITLE OF INVENTION: THERAPEUTIC MONOCLONAL ANTIBODIES TITLE OF INVENTION: NEUROTOXINS TILE OF INVENTION: NEUROTOXINS TILE OF INVENTION NUMBER: US/10/632,706 CURRENT APPLICATION NUMBER: US/10/632,706 CURRENT APPLICATION NUMBER: US 60/400,721 PRIOR PILLING DATE: 2003-08-01 PRIOR PILLING DATE: 2002-08-01 PRIOR APPLICATION NUMBER: US 60/400,721 PRIOR PILLING DATE: 1208-08-31 NUMBER OF SEQ ID NUMBER: US 09/144,806 PRIOR FILING DATE: 1998-08-31 NUMBER OF SEQ ID NOS: 278 SOFTWARE: Patentin version 3.2 EQ ID NO 126 LENGTH: 11 TYPE: PRT ORGANISM: Artificial	ALIGNMENTS	US-10-292-088-106 US-10-041-840-31 US-10-041-860-243 US-10-0465-383-42 US-10-465-383-42 US-10-465-383-19 US-10-364-743-109 US-10-364-743-110 US-10-364-743-110 US-10-452-593-109 US-10-452-593-109 US-10-452-593-110 US-10-452-593-110 US-10-452-593-110 US-10-452-593-110 US-10-452-593-109
THAT NEUTRALIZE BOTULINUM		Sequence 106, Applequence 31, Apple Sequence 243, Apple Sequence 42, Apple Sequence 53, Apple Sequence 53, Apple Sequence 10, Apple Sequence 16, Apple Sequence 110, Apple Sequence 140, Apple Sequence 140, Apple Sequence 140, Apple Sequence 34, Apple Sequence 61, Apple Sequ

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RESULT 2
US-10-632-706-127
(US-10-632-706-127)

§ Sequence 127, Application US/10632706

§ Publication No. US20040175385A1

GENERAL INFORMATION:

APPLICANT: MARKS, JAMES D.

APPLICANT: AMERSDOREER, PETER

TITLE OF INVENTION: THERAPEUTIC MONOCLONAL ANTIBODIES THAT NEUTRALIZE BOTULINUM

FILE REFERENCE: 4077-895120US

FULL REFERENCE: 4077-895120US

CURRENT APPLICATION NUMBER: US/10/632,706

CURRENT FILING DATE: 2003-08-01

PRIOR APPLICATION NUMBER: US 09/144,806
                                                                                                                                                                                                                                                                                                                                                                                                               ; FEATURE:
; OTHER INFORMATION: single chain antibody fragment
US-10-632-706-126
                                                                                                                                                                                                                                                                                  문
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                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local (
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                                                                                                                                                                                                                                                                                                                                                 11; Conservative
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                     0.0027;
thes 0;
                                                                                                                                                                                                                                                                                                                                                                                  DB 4; Length 11;
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Sequence 101, Application US/10364743
Publication No. US20040009178A1
GENERAL INFORMATION:
APPLICANT: Browdish, Katherine S.
APPLICANT: Frederickson, Shana
APPLICANT: Wild, Martha A.
APPLICANT: Wild, Martha A.
APPLICANT: No. US20040009178A1an, Mary Jean
APPLICANT: No. US20040009178A1an, Mary Jean
TITLE OF INVENTION: IMMUNOTHERAPEUTICS FOR BIODEFENSE
FILE REFERENCE: 84 (1087-73)
CURRENT APPLICATION NUMBER: US/10/364,743
CURRENT APPLICATION NUMBER: US/60/428,807
PRIOR FILING DATE: 2002-11-25
NUMBER OF SEQ ID NOS: 118
CORPMANDE: DETAILS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: PATENTIN Ver. 3.3
SEQ ID NO 116
LENGTH: 122
TYPE: PRT
ORGANISM: Homo sapiens
US-11-031-485-116
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; FEATURE:
; OTHER INFORMATION: single chain antibody fragment
US-10-632-706-127
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                                                                                                                                                                                                                                                                                                                                                                                                                    US-10-364-743-101
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Best Local Similarity
Matches 7; Conserv
SOFTWARE: PatentIn version 3.2
SEQ ID NO 101
LENGTH: 129
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 100.0%; Score 60; DB 4; Length 11; Best Local Similarity 100.0%; Pred. No. 0.0027; Matches 11; Conservative 0; Mismatches 0; Indels
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NUMBER OF SEQ ID NOS: 278
SOFTWARE: PatentIn version 3.2
SEQ ID NO 127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 116, Application US/11031485 Publication No. US20050232917A1
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APPLICANT: MOLLOY, BLIZABETH
APPLICANT: KELLERMANN, SIRID-AIMEE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 11
TYPB: PRT
ORGANISM: Artificial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: GREEN, LARRY L.
APPLICANT: HAAK-FRENDSCHO, MARY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                63.6%;
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Pred. No. 2.8;
3; Mismatches 1; Indels
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APPLICANT: Bowdish, Katherine S.
APPLICANT: Frederickson, Shana
APPLICANT: Wild, Martha A.
APPLICANT: Wild, Martha A.
APPLICANT: Molid, Martha A.
APPLICANT: Molidon Toshiaki
APPLICANT: Molidon Molidon Mary Jean
TITLE OF INVENTION: IMMUNOTHERAPEUTICS FOR BIODEFENSE
FILE REFERENCE: 84 (1087-73)
CURRENT APPLICATION NUMBER: US/10/364,743
CURRENT APPLICATION NUMBER: US 60/428,807
PRIOR APPLICATION WINDER: US 60/428,807
PRIOR FILING DATE: 2002-11-25
NUMBER OF SEQ ID NOS: 118
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; ORGANISM: huma
US-10-364-743-101
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Publication No. US20040009178A1
GENERAL INFORMATION:
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LENGTH: 129
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Matches 7; Conservative
SOFTWARE: PatentIn version 3.2 SEQ ID NO 15 LENGTH: 134
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APPLICANT: Bowdish, Katherine S.
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APPLICANT: Wild, Martha A.
APPLICANT: Wild, Martha A.
APPLICANT: Maryama, Toshiaki
APPLICANT: Nolan, Mary Jean
TITLE OF INVENTION: IMMUNOTHERAPEUTICS FOR BIODEFENSE
FILE REPERENCE: 98 CIP (1087-73 CIP)
CURRENT APPLICATION NUMBER: US/10/452,593
CURRENT FILING DATE: 2003-06-02
CURRENT FILING DATE: 2003-06-02
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PRIOR FILING DATE: 2003-02-11
PRIOR APPLICATION NUMBER: US 60/356,086
PRIOR FILING DATE: 2002-02-11
PRIOR APPLICATION NUMBER: US 60/376,408
PRIOR APPLICATION NUMBER: US 60/428,807
PRIOR APPLICATION NUMBER: US 60/428,807
PRIOR ETILING DATE: 2002-04-29
PRIOR ETILING DATE: 2002-11-26
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORGANISM: human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: PRT
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o. US20040258699A1
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Pred. No. 4.4;
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; ORGANISM: human US-10-364-743-15

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US-11-031-485-56

Sequence 56, Application US/11031485

Publication No. USCO050232917A1

GENERAL INFORMATION:

APPLICANT: HOLLOY, ELIZABETH
APPLICANT: WOLLOY, ELIZABETH
APPLICANT: KELLERMANN, SIRID-AIMEE
APPLICANT: KELLERMANN, SIRID-AIMEE
APPLICANT: HAAK-FRENDSCHO, MARY
APPLICANT: HAAK-FRENDSCHO, MARY
TITLE OP INVENTION ANTIBODIES TO MACCAM
FILE REFERENCE: ABX-PF6
CURRENT APPLICATION NUMBER: US/11/031,485
CURRENT APPLICATION NUMBER: 60/535,490
PRIOR APPLICATION NUMBER: 60/535,490
PRIOR TILING DATE: 2004-01-09
NUMBER OF SEQ ID NOS: 147
SOFTWARE: PACENTIN Ver. 3.3

SEQ ID NO 56
LENGTH: 468
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; ORGANISM: human
US-10-452-593-15
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Best Local Similarity 77.8
".--ches 7; Conservative
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Publication No. US20040258699A1
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ORGANISM: Homo sapiens
                                         ENGTH: 468
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Wild, Martha A.
Maruyama, Toshiaki
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77.8%;
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77.8%;
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Pred. No. 4.5;
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; ORGANISM: Homo sapiens
US-11-031-485-18
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                                                                             OTHER INFORMATION: single chain antibody fragment US-10-632-706-114
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US-10-632-706-114
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CURRENT APPLICATION NUMBER: US/11/031,485
CURRENT FILING DATE: 2005-01-07
PRIOR APPLICATION NUMBER: 60/535,490
PRIOR FILING DATE: 2004-01-09
NUMBER OF SEQ ID NOS: 147
SOFTWARE: PATENTIN VET. 3.3
SEQ ID NO 18
                                                                                                                           TITLE OF INVENTION: THERAPEUTIC MONOCLONAL ANTIBODIES THAT NEUTRALIZE BOTULINUM
TITLE OF INVENTION: NEUROTOXINS
FILE REFERENCE: 407T-895120US
CURRENT APPLICATION NUMBER: US(10/632,706
CURRENT FILING DATE: 2003-08-01
PRIOR APPLICATION NUMBER: US 60/400,721
PRIOR APPLICATION NUMBER: US 09/144,806
PRIOR FILING DATE: 2002-08-01
PRIOR APPLICATION NUMBER: US 09/144,806
PRIOR FILING DATE: 1998-08-31
NUMBER OF SEQ ID NOS: 278
SOFTWARE: Patentin version 3.2
SEQ ID NO 114
LENGTH: 13
TYPE: PRI
ORGANISM: Artificial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 114, Applic Publication No. US20 GENERAL INFORMATION:
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Best Local Similarity
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Best Local Similarity 77.8%;
Matches 7; Conservative
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Query Match 76.7%;
Best Local Similarity 70.0%;
Matches 7; Conservative
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APPLICANT: MOLLOY, BLIZABETH
APPLICANT: KELLERWANN, SIRID-AIMBE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: GREEN, LARRY L.
APPLICANT: HAAK-FRENDSCHO, MARY
TITLE OF INVENTION: ANTIBODIES TO MACCAM
                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: MARKS, JAMES I APPLICANT: AMERSDORFER,
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TYPE: PRT
                                                                                                                   FEATURE:
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  Score 46; DB 4; Length 13; Pred. No. 0.64; 2; Mismatches 1; Indels
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APPLICANT: Nixon, Andrew
APPLICANT: Hogan, Shannon
ITITLE OF INVENTION: PAPP-A LIGANDS
FILE REFERENCE: 10280-059001
CURRENT EAPPLICATION NUMBER: US/10/783,311
CURRENT FILING DATE: 2004-02-19
PRIOR APPLICATION NUMBER: US 60/448,515
PRIOR FILING DATE: 2003-02-19
INTERIOR PARTICING DATE: 394
SOFTWARE: PASCSQ for Windows Version 4.0
SEQ ID NO 135
RESULT 13
US-10-783-311-277
; Sequence 277, Application US/10783311
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US-10-783-311-135
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US-10-783-311-141
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US-10-783-311-135
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APPLICANT: Hogan, Shannon
TITLE OF INVENTION: PAPP-A LIGANDS
FILE REFERENCE: 10280-059001
CURRENT APPLICATION NUMBER: US/10/783,311
CURRENT FILING DATE: 2004-02-19
PRIOR APPLICATION NUMBER: US 60/448,515
PRIOR FILING DATE: 2003-02-19
PRIOR FILING DATE: 2003-02-19
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Publication No. US20050009136A1
GENERAL INFORMATION:
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Best Local Similarity
Matches 7; Conserv
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 141
LENGTH: 21
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Pred. No. 1;
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US-10-293-418-3068
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Sequence 3068, Application US/10293418
Publication No. US20030223996A1
GENERAL INFORMATION:
APPLICANT: Ruben et al.
TITLE OF INVENTION: Antibodies that Immunospecifically Bind BLyS
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APPLICANT: HOSAN, Andrew
APPLICANT: HOSAN, Shannon
TITLE OF INVENTION: PAPP-A LIGANDS
TITLE REFERENCE: 10280-059001
CURRENT APPLICATION NUMBER: US/10/783,311
CURRENT APPLICATION NUMBER: US-19
PRIOR APPLICATION NUMBER: US-19
PRIOR APPLICATION NUMBER: 003-02-19
PRIOR FILING DATE: 2003-02-19
NUMBER OF SEQ ID NOS: 394
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 277
LENGTH: 16
TYPE: PRI
DECANION: Attificial General
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US-10-783-311-277
                                                                                                                                                                            US-09-880-748-3068
                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: 60/212,210
PRIOR FILING DATE: 2000-06-15
PRIOR APPLICATION NUMBER: 60/240,816
PRIOR APPLICATION NUMBER: 60/240,816
PRIOR FILING DATE: 2000-10-17
PRIOR APPLICATION NUMBER: 60/276,248
PRIOR APPLICATION NUMBER: 60/277,379
PRIOR APPLICATION NUMBER: 60/277,379
PRIOR FILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: 60/293,499
PRIOR FILING DATE: 2001-05-25
NUMBER OF SEQ ID NOS: 3239
PRIOR FILING DATE: 2001-05-25
NUMBER OF SEQ ID NOS: 3239
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                                                                                                                                                                                                                                                               SOFTWARE: PatentIn Ver. 2.0 SEQ ID NO 3068
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Best Local Similarity 63.6%;
Matches 7; Conservative
                                                                                       Best Local Similarity Matches 7; Conserv
                                                                                                                                 Query Match
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CURRENT APPLICATION NUMBER: US/09/880,748
CURRENT FILING DATE: 2001-06-15
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TITLE OF INVENTION: Antibodies that Immunospecifically Bind BLyS
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ORGANISM: Homo sapiens
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Pred. No. 1.5;
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CURRENT PLING DATE: 2002-11-27

CURRENT FILING DATE: 2002-11-27

PRIOR APPLICATION NUMBER: 60/331,469

PRIOR APPLICATION NUMBER: 60/331,469

PRIOR APPLICATION NUMBER: 60/30,489

PRIOR PLILING DATE: 2001-06-15

PRIOR APPLICATION NUMBER: 60/23,499

PRIOR PLILING DATE: 2001-06-15

PRIOR PLILING DATE: 2001-06-15

PRIOR PLILING DATE: 2001-06-15

PRIOR PLILING DATE: 2001-06-15

PRIOR PLILING DATE: 2001-06-25

PRIOR PLILING DATE: 2001-06-25

PRIOR PLILING DATE: 2001-06-25

PRIOR APPLICATION NUMBER: 60/277,379

PRIOR PLILING DATE: 2000-01-3-16

PRIOR APPLICATION NUMBER: 60/276,248

PRIOR APPLICATION NUMBER: 60/276,248

PRIOR APPLICATION NUMBER: 60/276,248

PRIOR APPLICATION NUMBER: 60/210,210

PRIOR APPLICATION NUM
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Result
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RESULT 2 US-11-144-248-8 US-11-144-248-8 ; Sequence 8, Application US/11144248 ; Publication No. US20050244408A1

GENERAL INFORMATION:

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APPLICANT: MOYER, James D.
APPLICANT: MOYER, James D.
APPLICANT: Gorvalan, Jose R.
APPLICANT: Gallo, Michael
ITILE OF INVENTION: ANTIBODIES TO INSULIN-LIKE GROWTH FACTOR I RECEPTOR
FILE REFERENCE: ABX-PF2
CURRENT APPLICATION NUMBER: US/11/144,248
CURRENT APPLICATION NUMBER: US/10/038,591
PRIOR APPLICATION NUMBER: US/10/038,591
PRIOR APPLICATION NUMBER: 60/259,927
PRIOR FILING DATE: 2002-01-04
PRIOR APPLICATION UNUMBER: 60/259,927
PRIOR FILING DATE: 2001-01-05
NUMBER OF SEQ ID NOS: 60
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 8
LENGTH: 124
TYPE: PRT
ORGANISM: Homo sapiens
US-11-144-248-8
RESULT 4
US-11-144-248-4
; Sequence 4, Application US/11144248
; Publication No. US20050244408A1
; GENERAL INFORMATION:
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Publication No. US20050244408A1
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Cohen, Bruce D.
APPLICANT: Beebe, Jean
APPLICANT: Miller, Penelope E.
APPLICANT: Moyer, James D.
APPLICANT: Moyer, James D.
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Best Local Similarity
Matches 6; Conserva
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-144-248-16
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Best Local Similarity
Watches 6; Conserve
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CURRENT APPLICATION NUMBER: US/11/144,248
CURRENT APPLICATION NUMBER: US/10/038,591
PRIOR APPLICATION NUMBER: US/10/038,591
PRIOR FILLING DATE: 2002-01-04
PRIOR APPLICATION NUMBER: 60/259,927
PRIOR FILLING DATE: 2001-01-05
NUMBER OF SEQ ID NOS: 60
                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn Ver. 2.1 SEQ ID NO 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Corvalan, Jose R.
APPLICANT: Gallo, Michael
TITLE OF INVENTION: ANTIBODIES TO INSULIN-LIKE GROWTH FACTOR I RECEPTOR
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APPLICANT: Beebe, Jean
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                                                                                                                                           107 YYYYGMDV 114
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llarity 75.0%;
Conservative
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Pred. No. 0.092;
2; Mismatches
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Pred. No. 0.092;
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APPLICANT: GAILO, MICHAel

APPLICANT: GAILO, MICHAel

FILE OF INVENTION: ANTIBODIES TO INSULIN-LIKE GROWTH FACTOR I RECEPTOR

FILE REFERENCE: ABX-FF2

CURRENT APPLICATION NUMBER: US/11/144,248

CURRENT FILING DATE: 2005-06-02

PRIOR APPLICATION NUMBER: US/10/038,591

PRIOR APPLICATION NUMBER: US/10/038,591

PRIOR FILING DATE: 2002-01-04

PRIOR FILING DATE: 2001-01-05

PRIOR FILING DATE: 2001-01-05

NUMBER OF SEQ ID NOS: 60

SOFTWARE: Patentin Ver. 2.1

SEQ ID NO 4

LENGTH: 174

TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                            APPLICANT: Cohen, Bruce D.
APPLICANT: Heebe, Jean
APPLICANT: Miller, Penelope B.
APPLICANT: Miller, Penelope B.
APPLICANT: Moyer, James D.
APPLICANT: Corvalan, Jose R.
APPLICANT: Corvalan, Jose R.
APPLICANT: Gallo, Michael
ITILE OF INVENTION. ANTIBODIES TO INSULIN-LIKE GROWTH FACTOR I RECEPTOR
FILE REFERENCE: ABX-PF2
CURRENT FILING DATE: 2005-06-02
PRIOR APPLICATION NUMBER: US/11/144,248
CURRENT FILING DATE: 2002-01-04
PRIOR APPLICATION NUMBER: 60/259,927
PRIOR APPLICATION NUMBER: 60/259,927
PRIOR FILING DATE: 2001-01-05
NUMBER OF SEQ ID NOS: 60
SOFTWARE: Patentin Ver. 2.1
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; ORGANISM: Homo sapiens
US-11-144-248-45
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US-11-144-248-45
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RESULT 6
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Matches
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                                                                                                                                                                                                                                                                           SEQ ID NO 45
LENGTH: 470
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Matches 6; Conserv
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les 6; Conserv
                                                             126 YYYYGMDV 133
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Moyer, James D.
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Pred. No. 0.13;
2; Mismatches
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US-11-144-248-46
, Sequence 46, Application US/11144248
; Publication No. US20050244408A1
; GENERAL INFORMATION:
; APPLICANT: Cohen, Bruce D.

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CURRENT APPLICATION NUMBER: US/11/144,248
CURRENT FILING DATE: 2005-06-02
PRIOR APPLICATION NUMBER: US/10/038,591
PRIOR FILING DATE: 2002-01-04
PRIOR APPLICATION NUMBER: 60/259,927
PRIOR FILING DATE: 2001-01-05
NUMBER OF SEQ ID NOS: 60
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; TYPE: PRT; ORGANISM: Homo sapiens US-11-144-248-46
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US-11-144-248-49
RESULT 4
US-11-14-248-50
US-11-144-248-50
| Sequence 50, Application US/11144248
| Publication No. US20050244408A1
| GENERAL INFORMATION:
| APPLICANT: Chen, Bruce D.
| APPLICANT: Beebe, Jean
                                                                                                                                                                                                                                                                                                                                         ; LENGTH: 470
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-144-248-49
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SEQ ID NO 46
LENGTH: 470
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APPLICANT: Gallo, Michael
TITLE OP INVENTION: ANTIBODIES TO INSULIN-LIKE GROWTH FACTOR I RECEPTOR
FILE REFERENCE: ABX-PF2
CURRENT APPLICATION NUMBER: US/11/144,248
CURRENT FILING DATE: 2005-06-02
PRIOR APPLICATION NUMBER: US/10/038,591
PRIOR PILING DATE: 2002-01-04
PRIOR APPLICATION NUMBER: 05/259,927
PRIOR APPLICATION NUMBER: 60/259,927
PRIOR APPLICATION NUMBER: MADERIA SECTION APPLICATION PRIOR PILING DATE: 2001-01-05
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APPLICANT:
APPLICANT:
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APPLICANT: Beebe, Jean
APPLICANT: Miller, Penelope E.
APPLICANT: Moyer, James D.
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APPLICANT: Gallo, Michael
TITLE OF INVENTION: ANTIBODIES TO INSULIN-LIKE GROWTH FACTOR I RECEPTOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILE REFERENCE: ABX-PF2
                                                                                                                                                                                                                                                   70.0%;
Local Similarity 75.0%;
es 6; Conservation
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75.0%;
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Pred. No. 0.35;
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Pred. No. 0.35;
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; ORGANISM: Homo sapiens
US-11-144-248-50
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TYPE: PRT
ORGANISM: Homo sapiens
US-11-144-248-20
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US-11-144-248-20
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US-11-017-550-42
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                                                                          Sequence 42, Application US/11017550 Publication No. US20050250183A1 GENERAL INFORMATION:
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APPLICANT: Moyer, James D.
APPLICANT: Corvalan, Jose R.
APPLICANT: Golio, Michael
TITLE OF INVENTION: ANTIBODIES TO INSULIN-LIKE GROWTH FACTOR I RECEPTOR
FILE REFERENCE: ABX-PF2
CURRENT PELICATION NUMBER: US/11/144,248
CURRENT FILING DATE: 2005-06-02
PRIOR APPLICATION NUMBER: US/10/038,591
PRIOR PILING DATE: 2002-01-04
PRIOR PILING DATE: 2002-01-04
PRIOR FILING DATE: 2001-01-05
PRIOR FILING DATE: 2001-01-05
PRIOR FILING DATE: 2001-01-05
PRIOR PILING DATE: 2001-01-05
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APPLICANT: Moyer, James D.
APPLICANT: Moyer, Jose R.
APPLICANT: Gallo, Michael
TITLE OF INVENTION: ANTIBODIES TO INSULIN-LIKE GROWTH FACTOR I RECEPTOR
FILE REFERENCE: ABX-PP2
CURRENT APPLICATION NUMBER: US/11/144,248
CURRENT FILING DATE: 2005-06-02
PRIOR APPLICATION NUMBER: US/10/038,591
PRIOR FILING DATE: 2002-01-04
PRIOR FILING DATE: 2002-01-04
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PRIOR FILING DATE: 2001-01-05
NUMBER OF SEQ ID NOS: 60
SOFTWARE: Patentin Ver. 2.1
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APPLICANT: The Scripps Research Institute APPLICANT: Schultz, Peter G APPLICANT: Wang, Lei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Cohen, Bruce D. APPLICANT: Beebe, Jean
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Local Similarity 75.0%;
10s 6; Conservative
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                                                                                                                                                                                                                                                                                                                                                    Score 40; DB 7
Pred. No. 0.18;
5; Mismatches
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Anderson, John C

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/ ORGANISM: Methanococcus jannaschii
US-11-017-550-42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILE REFERENCE: 54-000120US
CURRENT APPLICATION NUMBER: US/11/017,550
CURRENT FILING DATE: 2004-12-17
PRIOR APPLICATION NUMBER: US/10/126,927
PRIOR FILING DATE: 2002-04-19
PRIOR APPLICATION NUMBER: US 60/285,030
PRIOR APPLILING DATE: 2001-04-19
PRIOR APPLICATION NUMBER: US 60/355,514
PRIOR FILING DATE: 2002-02-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 112, Applicat Publication No. US2005 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 79
SOFTWARE: PatentIn version 3.1
SEQ ID NO 42
LENTH: 306
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 60.0%;
Best Local Similarity 66.7%;
Matches 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Zhang, Zhiwen
TITLE OF INVENTION: In Vivo Incorporation of Unnatural Amino Acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  10-839-799-112
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
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                                                                                                                                                                                         COUNTRY: USA
ZIF: 20007-5109
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/839,799
FILING DATE: 06-May-2004
CLASSIFICATION: «Unknown»
PRIOR APPLICATION DATA:
                APPLICATION NUMBER: US/08/646,265
FILING DATE: 09-SEP-1996
APPLICATION NUMBER: WO PCT/JP94/01763
FILING DATE: 19-OCT-1994
APPLICATION NUMBER: JP 5-291078
FILING DATE: 19-NOV-1993
ATTORNEY/AGENT INFORMATION:
NAME: WEGNER, Harold C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: Foley & Lardner
STREET: 3000 K Street, N.W., Suite 500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: RESHAPED HUMAN ANTIBODY TO HUMAN MEDULLOBLASTOMA CELLS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: OHTOMO, TOShihiko
SATO, Koh
TSUCHIYA, Masayuki
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            158 TYHYAGVDV 166
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Meggers, Eric L
Mehl, Ryan A
Pastrnak, Miro
Santoro, Stephen W
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STATE: D.C
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REGISTRATION NUMBER: 25,258
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o. US20050249726A1
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Pred. No. 2.4;
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RESULT 13
US-10-721-763-25
[Sequence 25, Application US/10721763]
Publication No. US20050249729A1
[GENERAL INFORMATION:
APPLICANT: KIRIN BEER KABUSHIKI KAISHA
TITLE OF INVENTION: ANTI TRAIL-R ANTIBODY
FILE REFERENCE: PH-1573-PCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  á
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FILE REFERENCE: PH-1573-PCT
CURRENT APPLICATION NUMBER: US/10/721,763
CURRENT FILING DATE: 2003-11-26
PRIOR APPLICATION NUMBER: JP2001-150213
PRIOR FILING DATE: 2001-05-18
PRIOR PPLICATION NUMBER: JP2001-243040
PRIOR PILING DATE: 2001-05-09
PRIOR FILING DATE: 2001-08-09
PRIOR PPLICATION NUMBER: JP2001-314489
PRIOR PILING DATE: 2001-10-11
NUMBER OF SEQ ID NOS: 45
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            CURRENT APPLICATION NUMBER: US/10/721,763
CURRENT FILING DATE: 2003-11-26
PRIOR APPLICATION NUMBER: JP2001-150213
PRIOR FILING DATE: 2001-05-18
PRIOR APPLICATION NUMBER: JP2001-243040
PRIOR FILING DATE: 2001-08-09
PRIOR FILING DATE: 2001-08-09
PRIOR FILING DATE: 2001-10-11
PRIOR FILING DATE: 2001-10-11
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US-10-721-763-29
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQ ID NO 29
LENGTH: 145
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: KIRIN BEER KABUSHIKI KAISHA
NUMBER OF SEQ ID NOS: 45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEX: 904136
INFORMATION FOR SEQ ID NO: 112:
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MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 112:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
LENGTH: 87 amino acids
TYPE: amino acid
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TELEPHONE: (202)672-5300
TELEFAX: (202)672-5399
                                                                                                                                                                                                                                                                                                                                                                                          128 YYYGMDV 134
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imilarity 71.4%;
Conservative
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Pred. No.
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RESULT 15

US-11-022-562-211
; Sequence 211, Application US/11022562
; Publication No. US20050249742A1
; Publication No. US20050249742A1
; GENERAL INFORMATION:
    APPLICANT: Ruprecht, Ruth M.
; APPLICANT: Shisong, Jiang
; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR MODULATING
TITLE OF INVENTION: A CYTOTOXIC T LYMPHOCYTE IMMUNE RESPONSE
; FILE REFERENCE: DEW-043CN
; CURRENT APPLICATION NUMBER: US/11/022,562
; CURRENT FILING DATE: 2004-12-22
; PRIOR APPLICATION NUMBER: PCT/US03/20322
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Pastrnak, Miro
APPLICANT: Santoro, Stephen W
APPLICANT: Santoro, Stephen W
APPLICANT: Zhang, Zhiwen
TITLE OF INKENTION: In Vivo Incorporation of Unnatural Amino Acids
FILE REFERENCE: 54-00120US
CURRENT APPLICATION NUMBER: US/11/017,550
CURRENT FILING DATE: 2004-12-17
PRIOR APPLICATION NUMBER: US/10/126,927
PRIOR APPLICATION NUMBER: US 60/265,030
PRIOR FILING DATE: 2002-04-19
PRIOR APPLICATION NUMBER: US 60/265,030
PRIOR APPLICATION NUMBER: US 60/355,514
PRIOR APPLICATION NUMBER: US 60/355,514
PRIOR APPLICATION NUMBER: US 60/355,514
PRIOR FILING DATE: 2002-02-06
NUMBER OF SEQ ID NOS: 79
SOPTWARE: PatentIn version 3.1
SEQ ID NO 37
TENGTH: 306
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; TYPE: PRT
; ORGANISM: Methanococcus jannaschii
US-11-017-550-37
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Sequence 37, Application US/11017550
Publication No. US20050250183A1
GENERAL INFORMATION:
APPLICANT: The Scripps Research Institute
APPLICANT: Schultz, Peter G
APPLICANT: Wang, Lei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOPTWARE: PatentIn Ver. 2.1
SEQ ID NO 25
LENGTH: 154
TYPE: PRT
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Best Local
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10-721-763-25
                                                                                                                                                                                                                                                                                                                                                                                                                  Match 55.0%;
Local Similarity 62.5%;
Les 5; Conservative
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Meggers, Eric L
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Pred. No. 1.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 33; DB 7; Length 306; Pred. No. 8;
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Search completed: November 21, Job time: 0.301465 secs
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                                                                                                                                                                          ; TYPE: PRT; ORGANISM: Hepatitis C Virus US-11-022-562-211
                                                                                                                                                                                                                 SOFTWARE: PastSEQ for Windows Version 4.0
SEQ ID NO 211
LENGTH: 2280
                                                                                                                 Matches
                                                                                                                                             Query Match
                                                                                                                 Best Local Similarity 70.0
                                                                                                                                                                                                                                                               PRIOR FILING DATE: 2003-06-27
PRIOR APPLICATION NUMBER: 60/392718
PRIOR FILING DATE: 2002-06-27
NUMBER OF SEQ ID NOS: 340
                                                         1414 ÁVAÝÝRGLDV 1423
                                                                                      2 ATYYYFGLDV 11
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                  2005, 12:33:51
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GenCore version 5.1.6 Copyright (c) 1993 - 2005 Compugen Ltd.

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Result
No.
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Maximum
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Perfect score:
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protein search, using sw model
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Maximum Match 100%
Listing first 45 summaries
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geneseqp190s:*
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## ALIGNMENTS

RESULT 1 AAR97326 Monoclonal antibody; humanised; mouse; framework region; FR; CDR; complementary determining region; anti-darcinoembryonic antigen; CEA; diagnosis; imaging; therapy; immune response. WO9611013-A1 Humanised 15-OCT-1996 AAR97326 standard; peptide; 14 AA. 05-OCT-1994; 28-SEP-1995; 18-APR-1996. Homo sapiens. AAR97326; (IMMU-) IMMUNOMEDICS INC monoclonal (first entry) 94US-00318157. 95WO-US011964. antibody heavy chain framework region.

New humanised anti-CEA monoclonal CDRs, used for diagnosis, imaging antibody - having engrafted murine and therapy of CEA-producing cancers.

WPI; 1996-209653/21. Hansen HJ, Armour KL;

Claim 7; Page 39; 62pp; English.

New humanised monoclonal antibodies (MABs) comprising the complementary determining regions (CDRs) of a parental murine class III anticarcinoembryonic (CEA) MAB engrafted to the framework regions (PRs) of a heterologous antibody which can be derived from any species including human, retain the anti-CEA binding specificity of the parental murine MAB but are less immunogenic in a human subject than the parental MAB. The humanised antibodies can be used in diagnosis, imaging and therapy of CEA conjugates show improved therapeutic results, decreased immune responses and decreased immune-mediated adverse effects compared to the parent antibody. This sequence corresponds to the second framework region of the heavy chain of the humanised MAB. See AAR97313-97333

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RESULT 2
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ID AAO17
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DT 08-A
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                                                                                                                                                                                                                                                                         The present invention relates to a human or humanised antibody (Ab) which specifically binds to fibroblast activating protein alpha (PAPalpha). The antibodies are useful for preparing a composition for the treatment of cancer, and for imaging tumours associated with activated stromal fibroblasts, such as colorectal cancer, non-small-cell lung cancer, breast cancer, head and neck cancer, ovarian cancer, lung cancer, bladder cancer, pancreatic cancer and metastatic brain cancer, and diseases associated with the same, such as inflammation and wound healing. The present sequence is a peptide described in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Park J,
Schmidt
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New human humanized antibody that specifically binds to fibroblasts activating protein alpha, useful for treating cancer or tumor, and is imaging tumors associated with activated stromal fibroblasts, e.g. or breast cancer.
                                                                                                                                                                                                     Sequence 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Fig 6C; 109pp; English.
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11-SEP-2000; 2000GB-00022216.
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invention
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DB; AAL46555.
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herapy; cancer; wound healing; inflammation; cytostatic.
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                                                   HFR4;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Immunopolypeptide for diagnosis and treatment of human autoimmune disease, e.g., human rheumatoid arthritis, comprises a polypeptide that binds to human glucose-6-phosphate isomerase.
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             Synthetic.
                                                                                                                                                                                                                                                                                                                                       Sequence 14 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 6; Fig 4B; 47pp;
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(BURT/) BURTON D R.
(SCHA/) SCHALLER M.
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                                                   immunoglobulin molecule; heavy chain framework region; HFR1; HFR2; HFR3;
HFR4; tumour imaging; protein chips assay; light chain framework region;
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Pred. No. 3.2e-05;
0; Mismatches 0;
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Pischer R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention relates to an immunoglobulin molecule comprising of one or more heavy chain framework regions comprising HFR1, HFR2, HFR3 or HFR4 and one or more light chain framework regions comprising LFR1, LFR3 LFR3 or LFR4; and complementarity determining regions (CDRs) comprising CDR-H1, CDR-H2, CDR-H3 and/or CDRH1, CDR-L2 or CDR-L3. The immunoglobulin is useful in therapeutic or diagnostic assays comprising ELISA, phage display, tumour imaging or protein chips assays. Further, the immunoglobulin is useful in screening assays for detecting molecules that immunoglobulin is useful in screening assays for detecting molecules that
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New immunoglobulin molecule, useful in therapeutic or diagnostic assays comprising EDISA, phage display, tumor imaging or protein chips assay or in screening assays for detecting molecules that bind to the immunoglobulin molecule.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 14 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 7; SEQ ID NO 2; 198pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     N-PSDB; ADI20654, ADI20663, ADI20672.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    14-SEP-2001; 2001US-0318904P
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                                                                                                                                                                                                                                                                                              02-DEC-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              bind to the immunoglobulin molecule. The present sequence represents
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                                                                                                                                                                  Mue ep
                                                                                                                                                                                                           BONT/A neutralising epitope; anti-botulinum neurotoxin antibody; toxin neutralisation; botulinum neurotoxin poisoning; mouse; hea
                                                                                                                                                                                                                                                                 Mouse heavy chain anti-BoNT-antibody framework 2 segid 144.
                                                                                                                                                                                                                                                                                                                                                   ADR38742 standard; peptide;
                                        31-AUG-1998;
01-AUG-2002;
                                                                                 01-AUG-2003; 2003US-00632706.
                                                                                                                                       US2004175385-A1
                                                                                                                                                                                            framework 2.
                                                                                                                                                                                                                                     antibacterial; antibody; botulinum neurotoxin type A; BoNT/A;
              (REGC ) UNIV CALIFORNIA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                             (first entry)
                                        98US-00144886.
2002US-0400721P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%;
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                                                                                                                                                                                                                                                                                                                                                      14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 81; DB 7; Le
; Pred. No. 3.2e-05;
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Best Local &
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                                           Similarity
WVRQAPGKGLEWVA 14
                WVRQAPGKGLEWVA 14
                                 Conservative
                                          100.0%;
                                  0;
                                           Score 81; DB 8;
Pred. No. 3.2e-05;
                                    Mismatches
                                                  Length 14
                                    Indels
                                    0
                                    Gaps
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RESULT 7
ADR38738
ID ADR3
XX ADR3
XX ADR3
XX MOUB
XX MOUB
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XX BONLS
XW ETAN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CC clone where (I) binds to and neutralises botulinum neurotoxin type A specific CC clone where (I) binds to and neutralises botulinum neurotoxin type A (BoNT/A). An isolated antibody (I) that specifically binds to an epitope cCC specifically bound by an antibody expressed by a clone chosen from clone c25, C25, C39, IC6, 3D12, B4, IF3, huC25, Arl, Ar2, WRI(V), WRI(T), 3-1, CC 3-8, 3-10 and ING1, where (I) binds to and neutralizes botulinum cCC emprising BONT/A neutralising epitope having an epitope that is specifically bound by an antibody expressed by clones as mentioned in (I) producing (I); and a composition (III) comprising several anti-cc different epitope of a botulinum neurotoxin antibodies, where each antibody is specific for a clifferent epitope of a botulinum neutralisation than the single antibodies in surplus. The following are disclosed; a pharmaceutical composition (C comprising (I); and a kit comprising (I) (I) is useful for neutralising a botulinum neurotoxin which (I) in surplus, where each of (I) is specific for a different epitope of the botulinum neurotoxin which (I) in surplus, where each of (I) is specific for a different epitope of the botulinum neurotoxin than the composition or fantibodies in surplus. (I) is useful for diagnosing the botulism correction of antibodies associated with botulinum neurotoxin than the cc single antibodies in surplus. (I) is useful for diagnosing the botulism correction or diagnosis of botulism. This is the amino acid sequence of mouse heavy chain anti-BoNT-antibody framework region 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local 9
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                                                                                                                           US2004175385-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Marks JD,
    31-AUG-1998;
                                        01-AUG-2003; 2003US-00632706.
                                                                                                                                                                                                                        antibacterial; antibody; botulinum neurotoxin type A; BoNT/A;
BoNT/A neutralising epitope; anti-botulinum neurotoxin antibody;
toxin neutralisation; botulinum neurotoxin poisoning; mouse; heavy chain;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 14 AA;
                                                                                  09-SEP-2004.
                                                                                                                                                                                                                                                                                                             Mouse heavy chain anti-BoNT-antibody framework 2 segid 140
                                                                                                                                                                                                                                                                                                                                                   02-DEC-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                    ADR38738 standard; peptide; 14 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention describes an isolated antibody (I) that specifically binds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 4, SEQ ID NO 156; 110pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    14; Conservative
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    98US-00144886
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%;
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Pred. No.
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The invention describes an isolated antibody (I) that specifically binds compensed by a specifically bound by an antibody expressed by a specific clone where (I) binds to and neutralises botulinum neurotoxin type A (CC (BoNT/A). An isolated antibody (I) that specifically binds to an epitope compensed by a specifically bound by an antibody expressed by a clone chosen from clone (S25, C25, C39, IC6, 3D12, B4, 1F3, huC25, Ar1, Ar2, WR1(V), WR1(T), 3-1, CC 3-8, 3-10 and ING1, where (I) binds to and neutralizes botulinum compensation (I) binds to an epitope that is specifically bound by an antibody expressed by a clone chosen from clone comprising BoNT/A neutralising epitope having an epitope that is specifically bound by an antibody expressed by clones as mentioned in (I) producing (I); and a composition (II) comprising several anticc of insurplus and polypeptide (II) comprising several anticc composition antibodies, where each antibody is specific for a different epitope of a botulinum neurotoxin and the combination of comprising (I); and a kit comprising (I). (I) is useful for neutralising CC subject of the comprising (I) and a kit comprising (I). (I) is useful for neutralising CC subject of a different epitope of the botulinum neurotoxin which (I) is specific for a different epitope of the botulinum neurotoxin than the composition of antibodies shows greater toxin neutralisation than the composition of antibodies shows greater toxin neutralisation than the composition of antibodies shows greater toxin neutralisation than the composition of antibodies shows greater toxin neutralisation than the composition of antibodies shows greater toxin neutralisation than the composition of antibodies shows greater toxin neutralisation than the composition of antibodies shows greater toxin neutralisation than the composition of antibodies shows greater toxin neutralisation than the composition of antibodies shows greater toxin neutralisation than the composition of the botulinum neutralisation of the botulinum courotoxin whic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New isolated antibody that neutralizes botulinum neurotoxin type A, useful for diagnosing botulism or for treating pathologies associated
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  14;
                            Similarity
  Conservative
                            100.0%;
  0
Score 81; DB Pred. No. 3.2 0; Mismatches
  . 3.2e-05;
                                                         DB 8; Length 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     antibacterial; antibody; botulinum neurotoxin type A; BoNT/A;
BoNT/A neutralising epitope; anti-botulinum neurotoxin antibody;
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01-AUG-2003; 2003US-00632706.

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ADR38746
ID ADR38746
ID ADR3748
AC ADR3
XX ADR3
XX ADR3
XX MOUS
DE MOUS
XX MOUS
XX MOUS
XX MUS2
PD 09-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention describes an isolated antibody (I) that specifically binds colone where (I) binds to and neutralises botulinum neurotoxin type A (BoNT/A). An isolated antibody (I), that specifically binds to an epitope specifically bound by an antibody expressed by a clone chosen from clone CC specifically bound by an antibody expressed by a clone chosen from clone CC 3-8, 3-10 and ING1, where (I) binds to and neutralizes botulinum cC neurotoxin type A (BoNT/A). Also described are: a polypeptide (II) cc comprising BoNT/A neutralising epitope having an epitope that is specifically bound by an antibody expressed by clones as mentioned in (I) i producing (I); and a composition (III) comprising several anticc different epitope of a botulinum neurotoxin antibodies, where each antibody is specific for a cdifferent epitope of a botulinum neurotoxin and the comprising (I); and a kit comprising (I) is useful for neutralising comprising (I); and a kit comprising (I) is useful for neutralising comprising (I); and a kit comprising a botulinum neurotoxin which comprising (I); and a kit comprising a botulinum neurotoxin and the composition of antibodies in surplus, where each of (I) is specific for a different epitope of the botulinum neurotoxin and the comprising (I); and a kit comprising a botulinum neurotoxin and the comprising (I); and a kit comprising (I) is useful for neutralising considered to the composition of antibodies in surplus. (I) in surplus, where each of (I) is specific for a different epitope of the botulinum neurotoxin than the comprising (I); and the comprising (I); and the comprising (I) is useful for an entralisation than the comprision of antibodies in surplus. (I) is supplus, where each of (I) is enabled antibodies in surplus associated with botulinum neurotoxin than the comprision of the botulism of the botulism of the comprision of mouse heavy chain anti-BoNT-antibody framework region 2.
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                  ADR38746 standard; peptide; 14 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 14 AA;
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01-AUG-2002;
                                                                                                                                           antibacterial; antibody; botulinum neurotoxin type A; BONT/A; BONT/A neutralising epitope; anti-botulinum neurotoxin antibody; toxin neuralisation; botulinum neurotoxin poisoning; mouse; heavy chain;
                                                                                                                                                                                                                                   Mouse heavy chain anti-BoNT-antibody framework 2 seqid 148
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2002US-0400721P.
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Pred. No. 3.2e-05;
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The invention describes an isolated antibody (I) that specifically binds CC to an epitope specifically bound by an antibody expressed by a specific CC clone where (I) binds to and neutralises botulinum neurotoxin type A CC (BoNT/A). An isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody (I) that specifically binds to an epitope CC 3-8, 3-10 and ING1, where (I) binds to and neutralizes botulinum CC 3-8, 3-10 and ING1, where (I) binds to and neutralizes botulinum CC comprising BoNT/A neutralising epitope Having an epitope that is CC specifically bound by an antibody expressed by clones as mentioned in (I) producing (I); and a composition (III) comprising several anticce different epitope of a botulinum neurotoxin antibody shows greater toxin neutralisation than the single antibodies shows greater toxin neutralisation than the single antibodies in surplus. The following are disclosed; a pharmaceutical composition (CC comprising (I); and a kit comprising (I) is useful for neutralising EONT/A antibody and for neutralising a botulinum neurotoxin which (CC involves contacting neurotoxin with (I) in surplus, where each of (I) is specific for a different epitope of the botulinum neurotoxin and the compination of antibodies shows greater toxin neutralisation than the CC or for treating pathologies associated with botulinum neurotoxin than the consideration of antibodies in surplus. (I) is useful for diagnosing the botulism (CC enables rapid detection or diagnosis of botulism. This is the amino acid sequence of mouse heavy chain anti-BoNT-antibody framework region 2.
                                                                                                                                                                                                                                                                                                               RESULT 10
ADR38766
                                                                                                                                                                                                                                                                                                                                                                                                                                S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New isolated antibody that neutralizes botulinum neurotoxin type A, useful for diagnosing botulism or for treating pathologies associated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       31-AUG-1998; 98US-00144886
01-AUG-2002; 2002US-0400721P
                                                                                            antibacterial; antibody; botulinum neurotoxin type A; BONT/A; BONT/A neutralising epitope; anti-botulinum neurotoxin antibody; toxin neutralisation; botulinum neurotoxin poisoning; mouse; heavy chain;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 4; SEQ ID NO 148; 110pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2004-652009/63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Marks JD, Amersdorfer P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-AUG-2003; 2003US-00632706
                                                                                                                                                                            Mouse heavy chain anti-BoNT-antibody framework 2 segid 168.
                                                                                                                                                                                                                 02-DEC-2004
                                                                                                                                                                                                                                                                                           ADR38766 standard; peptide; 14 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 14 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          with botulinum neurotoxin poisoning.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (REGC ) UNIV CALIFORNIA.
                                                                                                                                                                                                                                                        ADR38766
                                                                           framework
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                    WVRQAPGKGLEWVA 14
                                                                                                                                                                                                                                                                                                                                                                                               WVRQAPGKGLEWVA 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Score 81;
100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3.2e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0,
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RESULT 11
AD738758
ID AD738
XX
XX
AC AD738
XX
O2-DE
DT 02-DE
XX
MOUSE
XX
BONT,
KW BONT,
KW BONT,
KW COX11
KW Grame
XX
W Sown
XX
W Sown
XX
W Sown
XX
W Sown
XX
W Sown
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W Sown
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W Sown
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W Sown
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CT The invention describes an isolated antibody (I) that specifically binds CC clone where (I) binds to and neutralises botulinum neurotoxin type A CC (BNNT/A). An isolated antibody (I) that specifically binds to an epitope Specifically bound by an antibody expressed by a specific CC (BNNT/A). An isolated antibody (I) that specifically binds to an epitope CC specifically bound by an antibody expressed by a clone chosen from clone CC comprising BoNT/A neutralising epitope having an epitope that is CC specifically bound by an antibody expressed by clones as mentioned in (I); producing (I); and a composition (II) comprising several anti-CC producing (I); and a composition (II) comprising several anti-CC different epitope of a botulinum neurotoxin antibodies, where each antibody is specific for a cC comprising (I); and a composition neutralisation than the single antibodies shows greater toxin neutralisation than the single antibodies comprising (I); and a kit comprising (I) is useful for neutralising a botulinum neurotoxin which (I) is specific for a different epitope of the botulinum neurotoxin and the combination of antibodies shows greater toxin neutralisation than the specific for a different epitope of the botulinum neurotoxin and the combination of antibodies shows greater toxin neutralisation than the CC single antibodies in surplus. (I) is useful for diagnosing the botulism cC or or to treating pathologies associated with botulinum neurotoxin botulism cC or ables rapid detection or diagnosis of botulism. This is the amino acid cequence of mouse heavy chain anti-BoNT-antibody framework region 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           useful for diagnosing with botulinum neurotoxin poisoning.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        31-AUG-1998; 98US-00144886.
01-AUG-2002; 2002US-0400721P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    01-AUG-2003; 2003US-00632706
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               09-SEP-2004
                                                                      antibacterial; antibody; botulinum neurotoxin type A; BONT/A;
BONT/A neutralising epitope; anti-botulinum neurotoxin antibody;
toxin neutralisation; botulinum neurotoxin poisoning; mouse; hea
                                                                                                                                                        Mouse heavy
                                                                                                                                                                                                 02-DEC-2004
                                                                                                                                                                                                                                                                                   ADR38758 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 14 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New isolated antibody that neutralizes botulinum neurotoxin type A, useful for diagnosing botulism or for treating pathologies associated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Marks
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (REGC ) UNIV CALIFORNIA.
                                                   framework 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity 100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                            WVRQAPGKGLEWVA 14
                                                                                                                                                                                                                                                                                                                                                                                     WVRQAPGKGLEWVA 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQ ID NO 168; 110pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Amersdorfer P;
                                                                                                                                                      chain anti-BoNT-antibody framework 2 seqid 160
                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                peptide; 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 81; DB 8; 1
Pred. No. 3.2e-05;
; Mismatches 0;
                                                                         poisoning; mouse; heavy chain;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
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CXXXXXXXXXXXX
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                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                         Query Match
          antibacterial; antibody; botulinum neurotoxin type A; BoNT/A; BoNT/A neutralising epitope; anti-botulinum neurotoxin antibody; toxin neutralisation; botulinum neurotoxin poisoning; mouse; hea
                                                                                                                                            ADR38750 standard; peptide; 14 AA.
                                                                Mouse
                                                                                                                   ADR38750;
                                                                                                                                                                                                                                      1 WVRQAPGKGLEWVA 14
                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                Conservative
                                                                chain
                                                                                         (first entry)
                                                                anti-BoNT-antibody framework 2 seqid
                                                                                                                                                                                                                                                                  100.0%; Score 81; DB 8;
100.0%; Pred. No. 3.2e-05;
tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                          Length 14;
               mouse; heavy chain;
                                                                                                                                                                                                                                                                   0
                                                                                                                                                                                                                                                                   Gaps
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Mus

US2004175385-A1

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RESULT 13
ADW96736
ID ADW96
XX
AC ADW96
XX
AC ADW96
XX
XX
DT 21-AF
XX
XX
MONOC
KW MONOC
KW Glion
Monoclonal antibody; diagnosis; therapy; cancer; tumor; carcinoma; glioma; neoplasm; cytostatic; epidermal growth factor receptor;
                                                                                                                                                                 ADW96736 standard; peptide; 14 AA
                                                      Anti-EGFRvIII antibody 13.1.2 VH peptide #3.
                                                                                                                                ADW96736,
                                                                                           (first entry)
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RESULT 14
ADW96723
ID ADW96
XX
AC ADW96
XX
AC ADW96
XX
DT 21-AF

ADW96723 standard;

peptide; 14 AA.

21-APR-2005 ADW96723;

(first entry)

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WVRQAPGKGLEWVA 14

Gaps

0,

The invention describes an isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a specific clone where (I) binds to and neutralises botulinum neurotoxin type A (BONT/A). An isolated antibody (I) that specifically binds to an epitope New isolated antibody that neutralizes botulinum neurotoxin type A, useful for diagnosing botulism or for treating pathologies associated 31-AUG-1998; 01-AUG-2003; 2003US-00632706. Example 4; SEQ ID NO 152; 110pp; English (REGC ) UNIV CALIFORNIA. 01-AUG-2002; 2002US-0400721P. botulinum neurotoxin poisoning. Amersdorfer P; 98US-00144886.

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CC (BONT/A). An isolated antibody (I) that specifically binds to an epitope CC (BONT/A). An isolated antibody expressed by a clone chosen from clone CC specifically bound by an antibody expressed by a clone chosen from clone CC specifically bound by an antibody expressed by a clone chosen from clone CC comprising BONT/A neutralising pitope having an epitope that is specifically bound by an antibody expressed by clones as mentioned in (I) comprising bound by an antibody expressed by clones as mentioned in (I) producing (I); and a composition (III) comprising several anticce different epitope of a botulinum neurotoxin antibody is specific for a CC different epitope of a botulinum neurotoxin and the combination of comprising (I); and a kit comprising (I). (I) is useful for neutralising comprising (I); and a kit comprising (I). (I) is useful for neutralising comprising (I); and a kit comprising (I). (I) is useful for neutralising BONT/A antibody and for neutralising a botulinum neurotoxin which (I) is comprising (I); and a kit comprising (I). (I) is useful for neutralising a botulinum neurotoxin and the combination of antibodies shows greater toxin neutralisation than the comprising of antibodies shows greater toxin neutralisation than the combination of antibodies shows greater toxin neutralisation than the combination of antibodies associated with botulinum neurotoxin and the combination of antibodies associated with botulinum neurotoxin correction or diagnosis of botulism. This is the amino acid correction of diagnosis of botulism. This is the amino acid correction of diagnosis of botulism. This is the amino acid correction of diagnosis of botulism.
Best Loc
Matches
                                                                                                       Query Match
                                                                                                                                                                                                                                Sequence 14 AA;
100.0%; Score 81; DB 8; 3
Similarity 100.0%; Pred. No. 3.2e-05;
L4; Conservative 0; Mismatches 0;
                                                                                                             Length 14;
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밁
             S
             1 WVRQAPGKGLEWVA 14
WVRQAPGKGLEWVA 14
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Creceptor, particularly to the type III deletion mutant (SEFRVIII). Also cincluded are a hybridoma cell line producing the above antibody, or its cransformed cell comprising a gene encoding the above antibody. Also cell transformed cell comprising a gene encoding the above antibody, inhibiting cell comprising a gene encoding the antibody, inhibiting cell copolynucleotide molecule comprising a mudleotide sequence encoding a heavy considering that the expression of EGFRVIII, an isolated constitute (comprising a container, a domposition contained in the container, and a package insert or label indicating that the composition comprises the antibody cited above), an assay kit centre the detection of EGFRVIII in mammalian tissues or cells (to screen for lung, colon, gastric, renal, prostate or ovarian carcinomas, the comprising an antibody that binds the antibody with the antipoen, if present, a comprising an antibody that binds the antipoen protein and means for comprising an antibody that binds the antipoen protein and means to comprising an antibody that binds the antipoen protein and means to comprising an antibody that binds the antipoen protein and means to comprising an antibody that binds the antipoen protein and means to comprising an antibody that binds the antipoen protein and means to comprising an antibody that binds the antipoen protein and means to comprising an antibody ariants to EGFRVIII and killing a targeted condition that the composition and methods are useful for diagnosing, preventing or treating diseases associated with the expression of EGFRVIII, such as cancer, gliomas, tunors and carcinomas. The present sequence is a peptide derived from a VH region of an anti-EGFRVIII antibody of the invention consecution of the antibody structure.
  Matches
                    Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New human monoclonal antibodies directed against type III deletion mutants of epidermal growth factor receptor (EGFRvIII), useful for diagnosing, preventing or treating diseases associated with EGFRvIII expression, e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2005-142884/15.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Weber R, Feng X, Foord O, Green L, Gudas J, Rathanaswami P, Raya R, Yang XD, Corvalan J, Kang J, King CT, Klakamp SL, Su QJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 27-JUN-2003; 2003US-0483145P
26-NOV-2003; 2003US-0525570P
15-APR-2004; 2004US-0562453P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10-FEB-2005.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO2005012479-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         heavy chain variable region.
                                                                                     Sequence 14 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 17; SEQ ID NO 122; 207pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (ABGE-) ABGENIX INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               sapiens
14; Conservative
                       100.0%;
  .
..
                       Score 81; DB 9;
Pred. No. 3.2e-05;
    Mismatches
                                              Length 14;
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      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Liu Y;
Jia X;
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Anti-EGFRvIII antibody 131 VH peptide

#3

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ADW80308
ID ADW8
XX
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                                                          RESULT 15
                                                                                                                                                                                         Query Match
Best Local S
Matches 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New human monoclonal antibodies directed against type III deletion mutants of epidermal growth factor receptor (EGFRVIII), useful for diagnosing, preventing or treating diseases associated with EGFRVII expression, e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Weber R, Feng X, Foord O, Green L, Gudas Rathanaswami P, Raya R, Yang XD, Corvalan Kang J, King CT, Klakamp SL, Su QJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Monoclonal antibody; diagnosis; therapy; cancer; tumor; carcinoma; glioma; neoplasm; cytostatic; epidermal growth factor receptor; heavy chain variable region.
                     ADW80308
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             27-JUN-2003; 2003US-0483145P
26-NOV-2003; 2003US-0525570P
15-APR-2004; 2004US-0562453P
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                                                                                                                                                                                                                                                                 Sequence 14 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 16; SEQ ID NO 109; 207pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25-JUN-2004; 2004WO-US020564.
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                                                                                                                                                                                           14;
                                                                                                                                                                                                           Similarity
                                                                                                                                          WVRQAPGKGLEWVA 14
                   standard; peptide; 14
                                                                                                                                                                                           Conservative
                                                                                                                                                                                                         100.0%;
                                                                                                                                                                                           0
                                                                                                                                                                                                           Score 81;
Pred. No. :
                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                           3.2e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     رم رد
                                                                                                                                                                                                                            Length 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Keyt B, Foltz I,
                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Liu Y;
Jia X;
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                                                                                                                                                                                The invention relates to a novel method for killing a target cell. The comprises contacting the cell with an isolated antibody or its mutant (BGFRVIII), the antibody being conjugated to a therapautic agent, which is a toxin chosen from AEFP, MMAE, AURISTATIN B, DM-1 and ZAP, and where the antibody comprises a heavy chain amino acid sequence chosen from antibodies 13.1.2, 131, 170, 150, 095, 250, EGFR variants are caused by gene rearrangement accompanied by gene amplification. Eight major cocurring variants of EGFR are known. EGFRVIII, which is the most commonly cocurring variant of EGFR in human cancers, comprises a 267 as in-frame delation in the extracellular domain. The method of the invention commistrates cytostatic activity and may be useful for inhibiting cell constrated with the expression of EGFRVIII or for compliferation associated with the expression of EGFRVIII. As such the method may be utilized, in vivo, on a mammal e.g. human, suffering from an epithelial cell cancer such as lung, colon, gastric, renal, prostate, companied by the sum and the prostate, glioblastoma or ovarian carcinoma. The current sequence is that cof a human anti-epidermal growth factor receptor type III variant correction and the prostate of a human anti-epidermal growth factor receptor type III variant
                                                                  Query Match
Best Local S
Matches 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New isolated antibody that binds to epidermal growth factor receptor type III mutant EGFRVIII and being conjugated to therapeutic agent such as toxin, useful for inhibiting cell proliferation associated with expression of EGFRVIII.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      27-JUN-2003; 2003US-0483145P
26-NOV-2003; 2003US-0525570P
15-APR-2004; 2004US-0562453P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cell death; monoclonal antibody; toxin; cytostatic; cancer; neoplasm; lung tumor; colon tumor; stomach tumor; renal tumor; prostatic cancer; breast tumor; oyary tumor; epidermal growth factor receptor; EGFRVIII;
                                                                                                                                     Sequence 14 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 18; SEQ ID NO 122; 233pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Weber R, Feng
Rathanaswami P,
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                                1 WVRQAPGKGLEWVA 14
                                                                                    Similarity
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, Raya R, Yang XD, Corvalan
CT, Klakamp SL, Su QJ;
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Pred. No. 3.
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Search completed: November 21, Job time: 19.7436 secs

2005, 12:20:01

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    protein search,

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seq length: 2000000000
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 BLOSUM62
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Copyright (c) 1993 - 2005 Compugen Ltd.
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first 45 summaries
                   using sw
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C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Date: 07-Unn-1990 #sequence_revision 07-Jun-1990 #text_change 31-Dec-2004
C;Accession: PLO120
R,Bird, J.; Galill, N.; Link, M.; Stites, D.; Sklar, J.
J. Exp. Med. 168, 229-245, 1988
A;Title: Continuing rearrangement but absence of somatic hypermutation in in A;Reference number: PLO116; MUID:88286083; PMID:2840480
A;Accession: PLO120
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PL0120
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                                                                                                                                                                                                                                                                         Matches
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81       100.0       115       2       S57445       Ig heavy chain V-J         81       100.0       115       2       S36267       Ig heavy chain V-I         81       100.0       116       1       M3HUGL       Ig heavy chain V-I         81       100.0       117       1       G1HUWS       Ig heavy chain V-I         81       100.0       117       2       S17079       Ig heavy chain V-I         81       100.0       117       2       S78486       Ig heavy chain V-I         81       100.0       118       2       S11677       Ig heavy chain V-I         81       100.0       118       2       PH1660       Ig heavy chain V-I         81       100.0       118       2       S11116       Ig heavy chain V-I         81       100.0       119       2       S31110       Ig heavy chain V-I         81       100.0       119       1       G1HUNI       Ig heavy chain V-I         81       100.0       119       1       M3HUM       Ig heavy chain V-I         81       100.0       119       1       M3HUM       Ig heavy chain V-I         81       100.0       119       2	45	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30
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## ALIGNMENTS

in immunoglobulin

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RESULT 2
PH0873
R, Manhetmer-Lory, A.; Katz, J.B.; Pillinger, M.; Ghossein, C.; Smith, A.; Diamond, J. Xxp. Med. 174, 1639-1652, 1991
A;Title: Molecular characteristics of antibodies bearing an anti-DNA-associated idi A;Accession: PH0873
R, Manhetmer-Lory, A.; Katz, J.B.; Pillinger, M.; Ghossein, C.; Smith, A.; Diamond, J. Xxp. Med. 174, 1639-1652, 1991
A;Title: Molecular characteristics of antibodies bearing an anti-DNA-associated idi A;Reference number: PH0862; MUID:92078875; PMID:1660528
A;Accession: PH0873
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Cross-references: UNIPROT:QBWUK1; UNIPROT:Q9UL93; UNIPARC:UPI0000176A2C
A;Experimental source: B cells from patient TD with acute lymphoblastic leukemia,
A;Experimental source: B cells from patient TD with acute lymphoblastic leukemia,
A;Note: the sequence shows the V region (TD-Vo) from a nonproductive DNA rearrange
C;Superfamily: immunoglobulin homology
C;Superfamily: immunoglobulin homology
C;Keywords: acute lymphoblastic leukemia; heterotetramer; immunoglobulin
F;31-35/Region: complementarity-determining 1
F;49-65/Region: complementarity-determining 2
                                                    A;Cross-references: UNIPARC:UPI000176C02
C;Comment: This antibody is produced by Epstein-Barr virus-transformed B cell that bears C;Commently: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
C;Keywords: heterotetramer; immunoglobulin
F;14-96/Domain: immunoglobulin homology <1MM>
F;14-96/Domain: immunoglobulin homology <1MM>
F;49-66/Region: complementarity-determining 1
                                                                                                                                                                                                                                                               A; Molecule type: DNA
A; Residues: 1-96 < MAN>
Query Match
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Pred. No. 4
      Score 81;
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         Length
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Ig heavy chain V region (anti-DNA, III-3R) - human (fragment)
C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Date: 09-Oct-1992 #sequence_revision 09-Oct-1992 #text_change 09-Jul-2004
C;Accession: PH0872
R;Manheimer-Lory, A.; Katz, J.B.; Pillinger, M.; Ghossein, C.; Smith, A.; Diamond, B.
J. Exp. Med. 174, 1639-1652, 1991
A;Tille: Molecular characteristics of antibodies bearing an anti-DNA-associated idiotype
A;Reference number: PH0862; MUID:92078875; PMID:1660528
A;Reference number: PH0862; MUID:92078875; PMID:1660528
A;Accession: PH0872
A;Molecule type: DNA
A;Residues: 1-97 <many
A;Cross-references: UNIFROT:09ULB6; UNIPARC:UPI0000176C04
A;Cross-references: UNIFROT:09ULB6; UNIPARC:UPI0000176C04
A;Cross-references: UNIFROT:09ULB6; UNIPARC:UPI0000176C04
A;Cross-references: UNIFROT:09ULB6; UNIPARC:UPI0000176C04
A;Cross-references: UNIFROT:09ULB6; UNIPARC:UPI0000176C04
A;Cross-references: UNIFROT:09ULB6; UNIPARC:UPI0000176C04
A;Cross-references: UNIFROT:09ULB6; UNIPARC:UPI0000176C04
A;Cross-references: UNIFROT:09ULB6; UNIPARC:UPI0000176C04
A;Cross-references: UNIFROT:09ULB6; UNIPARC:UPI0000176C04
A;Cross-references: UNIFROT:09ULB6; UNIPARC:UPI0000176C04
A;Cross-references: UNIFROT:09ULB6; UNIPARC:UPI0000176C04
A;Cross-references: UNIFROT:09ULB6; UNIPARC:UPI0000176C04
A;Cross-references: UNIFROT:09ULB6; UNIPARC:UPI0000176C04
A;Cross-references: UNIFROT:09ULB6; UNIPARC:UPI0000176C04
A;Cross-references: UNIFROT:09ULB6; UNIPARC:UPI0000176C04
A;Cross-references: UNIFROT:09ULB6; UNIPARC:UPI0000176C04
A;Cross-references: UNIFROT:09ULB6; UNIPARC:UPI0000176C04
A;Cross-references: UNIFROT:09ULB6; UNIPARC:UPI0000176C04
A;Cross-references: UNIFROT:09ULB6; UNIPARC:UPI0000176C04
A;Cross-references: UNIFROT:09ULB6; UNIPARC:UPI0000176C04
A;Cross-references: UNIFROT:09ULB6; UNIPARC:UPI0000176C04
A;Cross-references: UNIFROT:09ULB6; UNIPARC:UPI0000176C04
A;Cross-references: UNIFROT:09ULB6; UNIPARC:UPI0000176C04
A;Cross-references: UNIFROT:09ULB6; UNIPARC:UPI0000176C04
A;Cross-references: UNIFROT:09ULB6; UNIPARC:UPI0000
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A;Residues: 1-97 <HAM'>
A;Residues: 1-97 <HAM'>
A;Residues: 1-97 <HAM'>
A;Cross-references: UNIPARC:UPI000011662B; EMBL:Z31384; NID:g472969;
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
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Cipate: 13-Jan-1995 #sequence_revision 13-Jan-1995 #text_change 24-May-2001
Cipate: 13-Jan-1995 #sequence_revision 13-Jan-1995 #text_change 24-May-2001
Cipaccession: 844115
Rihawkins, R.B.; Zhu, D.; Ovecka, M.; Winter, G.; Hamblin, T.J.; Stevenson, F.K.
submitted to the EMBL Data Library, March 1994
Bibbacription: Idiotypic vaccination against human B-cell lymphoma: rescue of variable A; Reference number: 844105
A; Accession: 844115
A; Accession: 844115
A; Status: preliminary
A; Accession: 844115
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                            Ig heavy chain V-III region (AW-Vx) - human (fragment) C;Species: Homo sapiens (man) C;Dete: 07-Jun-1990 #sequence revision 07-Jun-1990 #te C;Accession: PI0116; $26892
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      Link, M.; Stites,
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Pred. No.
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i Mismatches 0;
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Pred. No. 4.8e-06;
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                                                             07-Jun-1990 #text_change 23-Jul-1999
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                                                                                      Ig heavy chain V-III region (TD-Vr) - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 07-Jun-1990 #sequence_revision 07-Jun-1990 #text_change
C;Accession: PLO123; S26897
R;Bird, J.; Galili, N.; Link, M.; Stites, D.; Sklar, J.
J. Exp. Med. 168, 229-245, 1988
A;Title: Continuing rearrangement but absence of somatic hypermu
A;Reference number: PLO116; MUID:88286083; PMID:2840480
A;Accession: PLO123.
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S29543
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                                   A; Molecule type: mRNA
A; Residues: 1-98 <BIR>
      A;Cross-references: UNIPARC:UPI0000116413
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Matches 14
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Best Local S
Matches 14
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A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-98 - TOM-
A;Crose-references: UNIPARC:UPI0000031F3A; EMBL:Z12349; NID:g32918; PIDN:CAA78219.1;
A;Crose-references: UNIPARC:UPI0000031F3A; EMBL:Z12349; NID:g32918; PIDN:CAA78219.1;
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: acute lymphoblastic leukemia; heterotetramer; immunoglobulin
F;15-98/Domain: immunoglobulin homology <IMM-
F;31-35/Region: complementarity-determining 1
F;49-65/Region: complementarity-determining 2</pre>
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A,Title: The repertoire of human germline V(H) sequences reveals about fifty group A,Reference number: $26885; MUID:93021117; PMID:1404388
A;Accession: $26882
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A;Title: Continuing rearrangement but absence of somatic hypermutation A;Reference number: PL0116; MUID:88286083; PMID:2840480
A;Accession: PL0116
                                                                                                                                                                                                                        A;Cross-references: UNIPARC:UPI000011649C; EMBL:Z17389; NID:g32835; C;Superfamily: immunoglobulin V region; immunoglobulin homology C;Keywords: heterotetramer; immunoglobulin F;15-98/Domain: immunoglobulin homology <IMM>
                                                                                                                                                                                                                                                                                                                                               A; Accession: S29543
A; Molecule type: DNA
A; Residues: 1-98 < TOM>
                                                                                                                                                                                                                                                                                                                                                                                                                                 R;Tomlinson, M.; Walter, G.; Cook, G.P.; Winter, submitted to the EMBL Data Library, October 1992 A;Reference number: S29543
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A; Residues: 1-98 <BIR>
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C;Date: 07-Jan-1994 #sequence_revision 17-Nov-1995
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                                                                                                                                            Local Similarity
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                                                                                                                Conservative
                                                                                                             100.0%; Score 81; DB 2;
100.0%; Pred. No. 4.8e-06;
tive 0; Mismatches 0;
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49
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Pred. No. 4.8e-06;
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but absence of somatic hypermutation 88286083; PMID:2840480

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immunoglobulin

23-Jul-1999

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A;Experimental source: B cells from patient TD with acute lymphoblastic leukemia, ALL A;Note: the sequence shows the V region (TD-Vr) from a nonproductive DNA rearrangement R;Tomlinson, I.M.; Walter, G.; Marks, J.D.; Llewelyn, M.B.; Winter, G. J. Mol. Biol. 227, 776-798, 1992
A;Title: The repertoire of human germline V(H) sequences reveals about fifty groups of A;Reference number: $26885; MUID:93021117; PMID:1404388
A;Accession: $26897
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A;Molecule type: DNA
A;Residues: 1-98 <TOM>
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R;Tomlinson, I.M.; Walter, G.; Marks, J.D.; Llewelyn, M.B.; Winter, G.
J. Mol. Biol. 227, 776-798, 1992
A;Title: The repertoire of human germline V(H) sequences reveals about fifty groups of A;Reference number: $26885; MUID:93021117; PMID:1404388
A;Accession: $26888
Ig heavy chain V region (COS-8 / DP-46) - numan (Iragment, C;Species: Homo sapiens (man) C;Species: 07-Jan-1994 #text_change 23-Jul-1999 C;Date: 07-Jan-1994 #text_change 23-Jul-1999
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R;Tomlinson, M.; Walter, G.; Cook, G.P.; Winter, G.submitted to the EMBL Data Library, October 1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                             ;Cross-references: UNIPARC:UPI000002DD16; EMBL:Z12346; NID:g32912; ;Note: designated DP-46
;Note: designated DP-46
;Superfamily: immunoglobulin V region; immunoglobulin homology ;Reywords: heterotetramer; immunoglobulin ;15-98/Domain: immunoglobulin homology <IMM>
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;Residues: 1-98 <TO2>
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;Residues: 1-98 <TOM>
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ilarity 100.0%;
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Pred. No. 4.8e-06;
); Mismatches 0;
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Pred. No. 4.8e-06;
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A;Accession: S20765
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-99 <MOR>
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                                                                                                                                                                                                                                                                                                                              R;Garcia, K.C.; Desiderio, S.V.; Ronco, P.M.; Verroust, P.J.; Amzel, L.M. Science 257, 528-531, 1992
A;Title: Recognition of angiotensin II: antibodies at different levels of an A;Reference number: A42575; MUID:92342952; PMID:1636087
A;Accession: B42575
                                                                                                                                                                                                                                                                                                                                                                                                                                        Ig heavy chain V region (anti-angiotensin II) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 05-Dec-1998 #sequence_revision 05-Ded-1998 #text_change 21-Jan-2000
C;Accession: B42575
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B42575
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submitted to the EMBL Data
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A;Residues: 1-101 <CARA
A;Residues: 1-101 <CARA
A;Residues: 1-101 <CARA
A;CEOSE-references: UNIPARC:UPI00001153E9; GB;S40679; NID:g252098; PIDN:AAB22669.1;
A;Experimental source: BALB/c mice, myeloma dell line NS-1
A;Note: sequence extracted from NCB; backbond (NCBIN:109337, NCBIP:109338)
A;Note: sequence extracted from NCB; backbond (NCBIN:109337, NCBIP:109338)
C;Superfamily: immunoglobulin v region; immunoglobulin homology
C;Keywords: immunoglobulin
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A;Molecule type: mRNA
A;Residues: 1-100 <KAV>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ig heavy chain V region - mouse
C;Species: Mus musculus (house mouse)
C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 08-Sep-1997
                                 S
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A;Accession: S26462
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Matches 14
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14; Conserv
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WVRQAPGKGLEWVA 41
                                     WVRQAPGKGLEWVA 14
                                                                                                                                                                 immunoglobulin homology <IMM>
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                                                                             100.0%; Score 81; DB 2; ilarity 100.0%; Pred. No. 5e-06; Conservative 0; Mismatches 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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100.0%; Pred. No. 4.9
tive 0; Mismatches
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Pred. No.
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hes 0;
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Ig heavy chain V region (clone 5D11) - human (fragment)
C;Species: Homo mapiens (man)
C;Species: Homo mapiens (man)
C;Date: 24-Feb-1994 #sequence_revision 24-Feb-1994 #text_change 31-Dec-2004
C;Accession: PH1644
R;Hillson, J.L.; Karr, N.S.; Oppliger, I.R.; Mannik, M.; Sasso, E.H.
J. Exp. Med. 178, 331-336, 1993
A;Title: The structural basis of germline-encoded VH3 immunoglobulin binding to staphylc A;Reference number: PH1642; MUID:93301610; PMID:8315388
A;Accession: PH1644
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C;Species: Homo sapiens (man)
C;Date: 24-Feb-1994 #sequence_revision 24-Feb-1994 #text_change 31-Dec-2004
C;Accession: PH1646
R;Hillson, J.L.; Karr, N.S.; Oppliger, I.R.; Mannik, M.; Sasso, E.H.
J. Exp. Med. 178, 331-336, 1993
A;Title: The structural basis of germline-encoded VH3 immunoglobulin binding to staphylc A;Reference number: PH1642; MUID:93301610; PMID:8315388
A;Accession: PH1646
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C;Species: Homo sapiens (man)
C;Date: 24-Reb-1994 #sequence_revision 24-Reb-1994 #text_change 31-Dec-2004
C;Accession: PH1642
R;Hillson: J.L.; Karr, N.S.; Oppliger, I.R.; Mannik, M.; Sasso, E.H.
J. Exp. Med. 178, 331-336, 1993
A;Title: The structural basis of germline-encoded VH3 immunoglobulin binding to staphylc
A;Reference number: PH1642; MUID:93301610; PMID:8315388
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PH1646
                                                                                A;Residues: 1-109 <HIL>
A;Residues: 1-109 <HIL>
A;Cross-references: UniproT:Q9UL93; UNIPARC:UPI0000176B7A
C;Superfamily: immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
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C;Superfamily: immunoglobulin homologuin
C;Keywords: heterotetramer; immunoglobulin
F;7-90/Domain: immunoglobulin homology <IMM>
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A; Molecule type: mRNA
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                                                    immunoglobulin homology <IMM>
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100.0%; Pred. No. 5.4e-06;
100.0%;
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Ig heavy chain V region (clone 2E8) - human (fragment)
C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Date: 24-Feb-1994 #sequence_revision 24-Feb-1994 #text_change 16-Aug-1996
C;Accession: PH1655
R;Hillson, J.L.; Karr, N.S.; Oppliger, I.R.; Mannik, M.; Sasso, E.H.
J. Exp. Med. 178, 331-336, 1993
A;Title: The structural basis of germline-encoded VH3 immunoglobulin binding to staphyloc A;Reference number: PH1642; MUID:93301610; PMID:8315388
A;Accession: PH1655
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PH1655
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A;Residues: 1-110 <HIL>
A;Cross-references: UNIPARC:UPI0000176BDC
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Best Local Similarity
Matches 14; Conserva
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WVRQAPGKGLEWVA 41
                                            WVRQAPGKGLEWVA 14
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                                                                                              Conservative
                                                                                            100.0%; Score 81; DB 2;
100.0%; Pred. No. 5.4e-06;
tive 0; Mismatches 0;
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Maximum Match 100%
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  protein search, using sw model
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MEDIJIN=75059123; PubMed=4803843;

Watanabe S., Barnikol H.U., Horn J., Bertram J., Hilschmann N.;

"The primary structure of a monoclonal IgM-immunoglobulin

(macroglobulin Gal.), II: the amino acid sequence of the H-chain
type), subgroup H III. Architecture of the complete IgM-molecule.
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Bur. J. Immunol. 22:247-251(1992).
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GO; GO:0003576; C:extracellular region; NAS.
GO; GO:0003823; F:antigen binding; NAS.
GO; GO:0006955; P:immune response; NAS.
InterPro; IPR007110; Ig-like.
InterPro; IPR007110; Ig-like.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG LIKE; 1.
Direct protein sequencing; Immunoglobulin do:
Immunoglobulin V region.
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NON TER
SEQUENCE
                                                                                                                                                          MEDLINE=93301610; PubMed=8315388; DOI=10.1084/jem.178.1.331; Hillson J.L., Karr N.S., Oppliger I.R., Mannik M., Sasso E.H. "The structural basis of germline-encoded VH3 immunoglobulin to staphylococcal protein A.";
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MEDLINE=98277119; PubMed=9614934; DOI=10.1006/clin.1998.4531;
MEDLINE=98277119; PubMed=9614934; DOI=10.1006/clin.1998.4531;
Min X.. Liu B., Van der Merwe P.L., Kalis N.N., Berney S.M.,
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Q\u00f3UL93;
Q\u001-mAY-2000 (TrEMBLrel. 13,
01-mAY-2000 (TrEMBLrel. 13,
01-mAR-2004 (TrEMBLrel. 26,
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Hilschmann N.;
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01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Myosin-reactive immunoglobulin heavy chain variable
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[2]
PubMed=2840480;
Bird J., Galili
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SMR; Q9UL93; 1-116.

InterPro; IPR007110; Ig-14ke.

InterPro; IPR003596; Ig_v.

SMART; SM00406; IGv; 1.

PROSITE; P850835; IG_LIKE; 1.
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"Variable region sequences of five human immunoglobulin heavy chains of the VH3 subgroup: definitive identification of four heavy chain hypervariable regions.";

Proc. Natl. Acad. Sci. U.S.A. 71:845-848(1974).

-I- MISCELLANEOUS: This chain was isolated from an IgG1 myeloma
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21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence update)
10-MAY-2005 (Rel. 47, Last annotation update)
11 heavy Chain V-III region WAS.
12 heavy Chain W-III region WAS.
13 heavy Chain (Human).
14 homo sapiens (Human).
15 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
16 homomalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
                                                                                                                                       GO; GO:0005576; C:extracellular region; NAS. GO; GO:000382; F:antigen binding; NAS. GO; GO:0006955; F:immune response; NAS. InterPro; IPR007110; Ig-like.

InterPro; IPR003596; Ig_v.
SMART; SM00406; IGv; 1.

PROSITE; PS50835; IG_LIKE; 1.

Direct protein sequencing; Immunoglobulin domain; Immunoglobulin V region.
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PIR; PL0120; PL0120.
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-!- SIMILARITY: Contains 1 Ig-like (immunoglobulin-like) domain.
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"Rule of antibody structure. The primary structure of a monoclonal 1961 immunoglobulin (myeloma protein Nie), I: purification and characterization of the protein, the L- and H-chains, the cyanogen bromide cleavage products, and the disulfide bridges.";
Hoppe-Seyler's Z. Physiol. Chem. 357:1515-1540(1976).
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hoppe-Seyler's Z. Physiol. Chem. 357:1571-1604(1976).
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Eukaryota; Metazoa; C
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GO; GO:0003823; F:antigen binding; NAS.
GO; GO:0006955; P:immune response; NAS.
InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
SMART; SM00406; IGv; 1.
PROSITE; PS59835; IG_LIKE; 1.
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10-MAY-2005 (Rel. 47, Last annot
Ig heavy chain V-III region NIE.
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SMR; P01770; 1-119.
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HV3M_HUMAN
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                                   HV3N HUMAN STANDARD; PRT; 1 P01775; 21-JUL-1986 (Rel. 01, Created) 21-JUL-1986 (Rel. 01, Last sequence up 10-MAY-2005 (Rel. 47, Last annotation Ig heavy chain V-III region LAY. Homo sapiens (Human).
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SEQUENCE
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21-JUL-1986 (Rel. 01, Last sequence up
10-MAY-2005 (Rel. 47, Last annotation
Ig heavy chain V-III region POM.
Homo sapiens (Human).
Eukaryota, Metazoa, Chordata; Craniata
Mammalia; Eutheria; Euarchontoglires;
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SMR; PO1774; 1-119.
GO; GO:0005576; C:extracellular region;
GO; GO:0003823; F:antigen binding; NAS.
GO; GO:0006955; P:immune response; NAS.
InterPro; IPR003596; Ig_v.
InterPro; IPR003596; Ig_v.
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Proc. Natl. Acad. Sci. U.S.A. 71:4032-4036(
-!- MISCELLANEOUS: This chain was isolated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDLINE=75046755; PubMed=4139708;
Capra J.D., Kehoe J.M.;
Capra J.D., Kehoe J.M.;
of the heavy chain variable regions of two immunoglobulin M anti-gamma
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P01774;
               Eukaryota; Metazoa;
Mammalia; Eutheria;
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PROSITE; PS50835; IG
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HSSP; P01772; 2FB4.
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-i- SIMILARITY: Contains 1 Ig-like (immunoglobulin-like)
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Direct protein sequencing; Immunoglobulin domain;

Immunoglobulin V region.

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1 112 Ig-like.

OMAIN

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119 AA;
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12954 MW;
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               Chordata; Craniata; Vertebrata; Euteleostomi
Euarchontoglires; Primates; Catarrhini; Homi
                                                                                                                                                                                                                                                      100.0%; Score 81;
100.0%; Pred. No.
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N -> D (probably doming isolation).
/FTId=VAR_003966.
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No. 3.2e-05;
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                Hominidae;
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RESULT 9
HV3J_HUMA
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Proc. Natl. Acad. Sci. U.S.A. 71:4034-1004.
Proc. Natl. Acad. Sci. U.S.A. 71:4034-1004.
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21-JUL-1986
10-MAY-2005
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P01771;
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PROSITE; PS50835; IG_LIKE; 1.
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MEDLINB=75046755; PubMed=4139708;
MEDLINB=75046755; PubMed=4139708;
Capra J.D., Kehoe J.M.;
"Structure of antibodies with shared idiotypy: the complete sequence
"Structure of antibodies with shared idiotypy: the complete sequence
"Structure of antibodies with shared idiotypy: the complete sequence
"Structure of antibodies with shared idiotypy: the complete sequence
                                                                This Swiss-Prot entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation - the European Bioinformatics Institute. There are no restrictions on its
                                                                                                                                                                                                                                           Chiu Y.-Y.H., Lopez de Castro J.A., Po
"Amino acid sequence of the VH region
cryolmmunoglobulin IgG Hil.",
Biochemistry 18:553-560(1979).
                                                                                                                                                                                                                                                                                                                 PROTEIN SEQUENCE.
MEDLINE=79124695; PubMed=420800;
Tonez de Castro J.A.,
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-i- SIMILARITY: Contains 1 Ig-like (immunoglobulin-like) domain.
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-I- SIMILARITY: Contains 1 Ig-like (immunoglobulin-like) domain.
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Mammalia; Eutheria;
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[1]
                                            use as
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-JUL-1986 (Rel. 01, Last sequence update)
-MAY-2005 (Rel. 47, Last annotation updat
heavy chain V-III region HIL.
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on of human myeloma
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RESULT 11
HV3K HUMAN
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AC P01772;
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EMBL; AF035030; AAD56286.1; -; mRNA.

HSSP; P01772; 2FB4.

SMR; Q9UL84; 1-122.

Ensembl; ENSG00000130076; Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Wu X., L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens (Human).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-CCT-2003 (TrEMBLrel. 25, Last annotation update)
Myosin-reactive immunoglobulin heavy chain variable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Q9UL84 HUMAN PRELIMINARY;
Q9UL84;
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GO; GO:0003823; F:antigen binding;
GO; GO:0006955; P:immune response;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDLINE=98277139; PubMed=9614934; DOI=10.1006/clin.1998.4531; WL X., Liu B., Van der Merwe P.L., Kalis N.N., Berney S.M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                      InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig-v.
SMART; SM00406; IGv; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                      fetus.";
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MEDLINE=83289131; PubMed=6884994;

MEDLINE=83289131; PubMed=6884994;

MEDLINE=83289131; PubMed=6884994;

MINITER OF COUNTY OF PAIR N., Hilschmann N.;

MINITER OF COUNTY OF PAIR N., Hilschmann N.;

BETTUCTURE OF CRYSTAllized monoclonal immunoglobulin IgG1 K

Hoppe-Seyler's Z. Physiol. Chem. 364:713-747(1983).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      pb; 2FB4; X-ray; H=2-126.

pb; 2FB4; X-ray; H=2-126.

pb; 2G2; X-ray; H=2-126.

G0; G0:000576; C:extracellular region; NAS.

G0; G0:000576; C:extracellular region; NAS.

G0; G0:000595; P:immune response; NAS.

InterPro; IPR007110; Ig-like.

InterPro; IPR007110; Ig-like.

InterPro; IPR007196; Ig-v.

SMART; SM00406; IGv; 1.

PROSITE; PS50835; IG-LIKE; 1.

3D-structure; Direct protein sequencing; Immunoglobulin domain;

Immunoglobulin V region; Pyrrolidone carboxylic acid.
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- SIMILARITY: Contains 1 Ig-like (immunoglobulin-like) domain.
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      13718 MW;
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Pyrrolidone carboxylic acid
        E4D71B52B16F8776 CRC64;
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Single-chain FV (Fragment).
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21-JUL-1986 (Rel. 01, Last sequence update)
10-MAY-2005 (Rel. 47, Last annotation update)
10 heavy chain V region IR2 precursor.
Rattus norvegicus (Rat).
Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalla; Butheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;
Maroidea; Muridae; Murinae; Rattus.
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SMR; P01805; 20-141.
Ensembl; ENSRNOG00000031589; Rattus
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MEDLINE=83064537; PubMed=6292865;
Hellman L., Petterson U., Engstroem A., Karlsson T.,
"Structure and evolution of the heavy chain from rat:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig v.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 1.
Immunoglobulin domain; Immunoglobulin V region; Signal.
Isignal
Ig heavy chain V region IR:
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20 142 Ig heavy chain V region IR:
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
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05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
Hypothetical protein DKFZp686I04196 (Fragment).
Name=DKFZp686I04196;
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MEDLINE=97362799; PubMed=9219263; DOI=10.1038/nbt0797-629;
KONTERMENT R.E., Wing M.G., Winter G.;
"Complement recruitment using bispecific diabodies.";
Nat. Biotechnol. 15:629-631(1997).
EMBL; Y13056; CAA73499.1; -; mRNA.
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	, ia &	,	187; N:	m as a		1815 L. PTEI	**	US-09-1 US-
	Score Pred. Mis		65 18733/464 N:	#1.0, #18,15	Suite	318157B L. APTED TYPE	TIG	00000000000000000000000000000000000000
	ore 81; ed. No. 2 Mismatche		164	7			ALIGNMENTS	20 00 00 00 00 00 00 00 00 00 00 00 00 0
	che		•	Version B	500	III	SILN	717-6 717-6 717-6 717-6 7117-6 7117-6 7117-6 7117-6 7117-6 7117-6 7117-6 7117-6 7117-6 7117-6
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	06;			#1.30		III ANTI-CEA ANTIBODIES		
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PATENT NO. 6676924

PATENT NO. 6676924

PATENT NO. 6676924

APPLICANT: HANSEN, Hans J.

ARMOUN, Kathryn L.

TITLE OF INVENTION: CDR-GRAFTED TYPE III ANTI-CEA HUMANIZED

MOUSE MONOCLONAL ANTIBODIES
                                                                                                                                                                                                                                                                  US-09-472-087-72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-09-253-794-33
                             GENERAL INFORMATION:
APPLICANT: HANGON, DOUGLAS C.
APPLICANT: NEVEU, MARK J.
APPLICANT: MUELLER, EILLEN B.
APPLICANT: HANKE, JEFFREY H.
APPLICANT: GILMAN, STEVEN C.
APPLICANT: DAVIS, C. GEOFFREY
APPLICANT: CORVALAN, JOSE R.
TITLE OF INVENTION: HUMAN MONOCLONAL ANTIBODIES TO CTLA-4
                                                                                                                                                                                                                       Patent No.
                                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
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Best Local
CURRENT APPLICATION NUMBER: US/09/472,087
                     FILE REFERENCE: ABX-PF1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEX: 904136
INFORMATION FOR SEQ ID NO: 33:
SEQUENCE CHARACTERISTICS:
LENGTH: 14 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION: AUDKNOWNS
PRIOR APPLICATION NUMBER: US 08/318,157
APPLICATION NUMBER: US 08/318,157
FILING DATE: 05-OCT-1994
ATTORNEY/AGENT INFORMATION:
NAME: SAXE, Bernhard D.
REGISTRATION NUMBER: 28,665
REFERENCE/DOCKET NUMBER: 18733/464
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA: D8/09/253,794 PILING DATE: 22-Feb-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: Foley & Lardner
STREET: 3000 K Street, N.W.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: linear MOLECULE TYPE: protein SEQUENCE DESCRIPTION: SEQ ID NO: 33:
                                                                                                                                                                                                                       72, Application US/09472087
o. 6682736
                                                                                                                                                                                                                                                                                                                                                                                                                              14; Conservative
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ZIP: 20007-5109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEPHONE: (202)672-5300
TELEFAX: (202)672-5399
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRANDEDNESS: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%; Score 81; DB 2; 100.0%; Pred. No. 2.9e-06;
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US-09-534-717-653
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                                                                                                                                                                                                                                                                                                                                                                                                  US-08-211-202-118
                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 5
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US-09-534-717-653
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GENERAL IN. 6914128

GENERAL INFORMATION:
APPLICANT: Jochen, Salfeld et al.
TITLE OF INVENTION: Human Antibodies That Bind Human IL-12 And Methods For Producing FILE REFERENCE: BBI-093CP
CURRENT APPLICATION NUMBER: US/09/534,717
CURRENT FILING DATE: 2000-03-24

EARLIER APPLICATION NUMBER: 60/126,603
EARLIER FILING DATE: March 25, 1999
NUMBER OF SEQ ID NOS: 675
CORTENAND SEQ ID NOS: 675
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Loc
Matches
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LENGTH: 95
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                                                                                                                                                                                                                                                                                                                                                                Sequence 118, Approximately Patent No. 55653
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PRIOR APPLICATION NUMBER: 60/113,647
PRIOR FILING DATE: 1998-12-23
NUMBER OF SEQ ID NOS: 147
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                            GENERAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT
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                                                                                                                                                                                                                                                                         APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                     STREET: 65.
STREET: Chicago
CITY: Chicago
Illinois
IISA
                                                                                                                                                                                                APPLICANT: BAIRR, Michael
APPLICANT: JESPERS, Laurent Stephane Anhe Therese
APPLICANT: WINTER, Gregory Paul
TITLE OF INVENTION: Production of chimeric antibodies -
TITLE OF INVENTION: combinatorial approach
NUMBER OF SEQUENCES: 14
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ENGTH: 89
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Local Similarity 100.0%;
                                                                                                                                                                ADDRESSEB: David W. Clough, Marshall O'Toole Gerstein Murray ADDRESSEB: Borun
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local
                                                                                                                                                                                                                                                                                                                                               INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         36 WVRQAPGKGLEWVA 49
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                                                                                                                                                                                                                                                                                                                                                                                   Application US/08211202
                                                                                                                                               6300 Sears Tower,
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illarity 100.0%;
Conservative 0;
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Pred. No. 2.4e-05;
                                                                                                                                               233 South Wacker Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 81;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
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US-08-211-202-118
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                                              COUNTRY: United States
ZIP: 20037-3202
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: HP 9000/700 Workstation
OPERATING SYSTEM: UNIX
SOFTWARE: In house
                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                  APPLICANT: PEDERSEN
APPLICANT: SEARLE,
APPLICANT: RES,
APPLICANT: ROGUSKA,
APPLICANT: GUILD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: David W. Clough
REGISTRATION NUMBER: 36,107
REFERENCE/DOCKET NUMBER: 28
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312-474-6300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: GB 9120252.3 FILING DATE: 23-SEP-1991 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: Patentin Release #1.0, Version #1.25 (EPO) CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/211,202 FILING DATE: 23-SEP-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
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APPLICATION NUMBER:
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/0:
FILING DATE: 09-SEP-1992
                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 5
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                               APPLICANT: GUILD,
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: PCT/GB92/00883
FILING DATE: 15-MAY-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: 24-MAR-1992
                                                                                                                                                                       CITY: Washington STATE: D.C.
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                                                                                                                                                                                                          STREET:
                                                                                                                                                                                                          ADDRESSEE: Sughrue, Mion, Zinn, Macpeak & Seas
STREET: 2100 Pensylvania Avenue, N.W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Match 100.0%;
Local Similarity 100.0%;
es 14; Conservative 0
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                                                                                                                                                                                                                                                                                                                                  ROGUSKA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            25-SEP-1991
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                                                                                                                                                                                                                                                                                                 Stephen M.J.
Anthony R.
Anthony R.
Michael A.
Braydon C.
Braydon C.
SURFACE RESIDUE VENEERING OF RODENT
                                                                                                                                                                                                                                                                                  ANTIBODIES
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                  US/07/942,245
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Pred. No. 2.5e-05;
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GENERAL INFORMATION:
APPLICANT: Foote, Jefferson
TITLS OF INVENTION: Super Humanized Antibodies
FILE REFERENCE: 501231.01
CURRENT APPLICATION NUMBER: US/10/194,975
CURRENT FILING DATE: 2002-10-10
PRIOR APPLICATION NUMBER: US 60/305,111
PRIOR APPLICATION NUMBER: US 60/305,111
PRIOR FILING DATE: 2001-07-12
NUMBER OF SEQ ID NOS: 122
SOFTWARE: PATENTIN Version 3.1
SEQ ID NO 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 7
US-10-194-975-15
; Sequence 15, Ap
; Patent No. 6881
                                                                                                                                                                                                                                                                                                        RESULT 8
US-10-194-975-23
; Sequence 23, Application US/10194975
; Patent No. 6881557
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-07-942-245-37
                                                    ; ORGANISM: Homo sapiens US-10-194-975-23
                                                                                                                                                    GENERAL INFORMATION:

APPLICANT: Poote, Jefferson

TITLE OF INVENTION: Super Humanized Antibodies

FILE REFERENCE: 50131.01

CURRENT APPLICATION NUMBER: US/10/194,975

CURRENT FILING DATE: 2002-10-10

PRIOR APPLICATION NUMBER: US 60/305,111

PRIOR FILING DATE: 2001-07-12

NUMBER OF SEQ ID NOS: 122

NUMBER OF SEQ ID NOS: 122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                     SOFTWARE: PatentIn version 3.1
SEQ ID NO 23
LENGTH: 98
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
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Query Match
Best Local Similarity
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 98
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION: 530
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 293-7060
TELEPAX: (202) 293-7860
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: amino acid
TOPOLOGY: linear
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Local Similarity 100.0%;
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Similarity 100.0%;
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   100.0%;
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Pred. No. 2.5e-05;
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Pred. No. 2.5e-05;
   Score 81; DB 2;
Pred. No. 2.5e-05;
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                   Length 98;
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FRUERIC NO. 100.101

[RENERAL INFORMATION: Gefferson
ITITLE OF INVENTION: Super Humanized Antibodies
FILE REFERENCE: 501231.01

[CURRENT APPLICATION NUMBER: US/10/194,975

[CURRENT FILING DATE: 2002-10-10
PRIOR APPLICATION NUMBER: US 60/305,111

PRIOR FILING DATE: 2001-07-12

NUMBER OF SEQ ID NOS: 122

SOPTWARE: Patentin version 3.1

SEQ ID NO 24

LENGTH: 98

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US-10-194-975-25
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US-10-194-975-24
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Sequence 26, Application US/10194975
PATENT NO. 6881557
GENERAL INFORMATION:
APPLICANT: Foote, Jefferson
TITLE OF INVENTION: Super Humanized
FILE REFERENCE: 501231.01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: POOTE, Jefferson
TITLE OF INVENTION: Super Humanized Antibodies
FILE REFERENCE: 501231.01
CURRENT APPLICATION NUMBER: US/10/194,975
CURRENT FILING DATE: 2002-10-10
PRIOR APPLICATION UNMBER: US 60/305,111
PRIOR FILING DATE: 2001-07-12
NUMBER OF SEQ ID NOS: 122
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Matches 14; Conservative 0; Mismatches n.
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SEQ ID NO 25
LENGTH: 98
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Patent No. 68815
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Best Local Similarity
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100.0%; Pred. No. 2.5e-05;
                                             Humanized Antibodies
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; ORGANISM: Homo sapiens US-09-534-717-624
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US-09-534-717-624
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US-10-194-975-26
                                                                                    ; ORGANISM: Homo sapiens
US-09-534-717-625
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Patent No. 6914128
GENERAL INFORMATION:
APPLICANT: Jochen, Salfeld et al.
TITLE OF INVENTION: Human Antibodies That Bind Human IL-12 And Methods For Producing FILE REFERENCE: BBI-093CP
CURRENT APPLICATION NUMBER: US/09/534,717
CURRENT FILING DATE: 2000-03-24
EARLIER APPLICATION NUMBER: 60/126,603
EARLIER FILING DATE: March 25, 1999
NUMBER OF SEQ ID NOS: 675
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 624
                                                                                                                                                             APPLICANT: Jochen, Salfeld et al.

APPLICANT: Jochen, Salfeld et al.

TITLE OF INVENTION: Human Antibodies That Bind Human IL-12 And Methods For Producing FILE REFERENCE: BBI-093CP
CURRENT APPLICATION UNMBER: US/09/534,717
CURRENT FILING DATE: 2000-03-24
EARLIER APPLICATION NUMBER: 60/126,603
EARLIER FILING DATE: March 25, 1999
NUMBER OF SEQ ID NOS: 675
SOFTWARE: Patentin Ver. 2.0

SOFTWARE: Patentin Ver. 2.0
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SEQ ID NO 26
LENGTH: 98
TYPE: PRT
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Query Match 100.0%; Score 81; DB 2; 1
Best Local Similarity 100.0%; Pred. No. 2.5e-05;
Matches 14; Conservative 0; Mismatches 0;
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Best Local Similarity 100.0%;
Matches 14; Conservative (
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Patent No. 6914128
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CURRENT FILING DATE: 2002-10-10
PRIOR APPLICATION NUMBER: US 60/305,111
PRIOR FILING DATE: 2001-07-12
NUMBER OF SEQ ID NOS: 122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 98
TYPE: PRT
                                                                                                                              TYPE: PRT
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Local Similarity 100.0%;
les 14; Conservative (
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Pred. No. 2.5e-05;
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Pred. No. 2.5e-05;
0; Mismatches 0;
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RESULT 15
US-09-534-717-627
; Sequence 627, Application US/09534717
; Sequence 627, Application US/09534717
; Patent No. 6914128
; GENERAL INFORMATION:
APPLICANT: Jochen, Salfeld et al.
APPLICANT: Jochen, Salfeld et al.
TITLE OF INVENTION: Human Antibodies That Bind Human IL-12 And Methods For Producing
FILE REFERENCE: BBI-093CP
CURRENT APPLICATION NUMBER: US/09/534,717
; CURRENT APPLICATION NUMBER: US/09/534,717
; CURRENT FILING DATE: APIC: 60/126,603
; EARLIER APPLICATION NUMBER: 60/126,603
; EARLIER FILING DATE: March 25, 1999
; NUMBER OF SEQ ID NOS: 675
; SOFTWARE: Patentin Ver. 2.0
; SEQ ID NO 627
; LENGTH: 98
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-534-717-627
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; CURRENT FILING DATE: 2000-03-24
; EARLIER APPLICATION NUMBER: 60/126,603
; EARLIER FILING DATE: March 25, 1999
; NUMBER OF SEQ ID NOS: 675
; SOFTWARE: Patentin Ver. 2.0
; SEQ ID NO 626
; LENGTH: 98
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-534-717-626
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Search completed: November 21, 2005, 12:07:38 Job time: 5.28205 secs
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APPLICANT: Jochen, Salfeld et al.

TITLE OF INVENTION: Human Antibodies That Bind Human IL-12 And Methods For Producing FILE REFERENCE: BBI-093CP
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                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                           y Match 100.0%; Score 81; DB 2; Length 98;
Local Similarity 100.0%; Pred. No. 2.5e-05;
hes 14; Conservative 0; Mismatches 0; Indels
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GenCore version 5.1.6 Copyright (c) 1993 - 2005 Compugen Ltd.

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Result
No.
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Maximum DB seq length: 200000000
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Perfect score:
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     Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score
     protein search, using
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query
Match
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81
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1867569 seqs, 417829326 residues
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1: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*

2: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

3: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

4: /cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*

5: /cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.pep:*

6: /cgn2_6/ptodata/1/pubpaa/US11_PUBCOMB.pep:*
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Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length
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                   US-09-253-794-33
US-09-828-708-64
US-10-632-706-144
US-10-632-706-154
US-10-632-706-154
US-10-632-706-156
US-10-632-706-156
US-10-632-706-160
US-10-632-706-164
US-10-632-706-164
US-10-632-706-164
US-10-933-068-202
US-10-923-068-202
US-10-923-068-202
US-10-923-068-202
US-10-877-773-1122
US-10-877-774-129
US-10-877-774-129
US-10-877-774-129
US-10-877-774-129
US-10-877-774-129
US-10-877-774-129
US-10-877-774-129
US-10-877-774-129
US-10-877-774-139
US-10-984-9608-337
US-10-923-068-334
US-10-923-068-334
US-10-923-068-336
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Sequence 33, Appl
Sequence 144, App
Sequence 144, App
Sequence 148, App
Sequence 156, App
Sequence 156, App
Sequence 160, App
Sequence 164, App
Sequence 168, App
Sequence 202, App
Sequence 202, App
Sequence 203, App
Sequence 204, App
Sequence 119, App
Sequence 122, App
Sequence 122, App
Sequence 123, App
Sequence 124, Appl
Sequence 127, App
Sequence 128, App
Sequence 129, App
Sequence 120, App
Sequence 121, Appl
Sequence 122, App
Sequence 123, App
Sequence 124, Appl
Sequence 334, Appl
Sequence 341, Appl
Sequence 364, Appl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Description
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US-09-253-794-33
   Query Match 100.0%; 9 Best Local Similarity 100.0%; F Matches 14; Conservative 0;
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ALIGNMENTS	US-10-194-975-23	US-10-194-975-15	US-09-948-939-21	US-09-948-939-15	US-10-884-830-653	US-10-363-204-91	US-10-363-204-89	US-09-910-483-45	US-09-910-483-39	US-10-776-649-72	US-10-612-497-72	US-10-842-011-45	US-10-788-625-90	US-10-230-880-123	US-10-230-880-122	US-10-230-880-121	US-10-230-880-120	US-10-230-880-119
	Sequence 23, Appl	Sequence 15, Appl	Sequence 21, Appl	15,	-			4.5	-	72,			90,	123,	122,	121,	120,	Sequence 119, App

## STRANDEDNESS: < Unknown> TOPOLOGY: linear MOLECULE TYPE: protein SEQUENCE DESCRIPTION: SEQ ID NO: US-09-253-794-33 Sequence 33, Application US/09253794 Patent No. US20020018750A1 GENERAL INFORMATION: TELEX: 904136 INFORMATION FOR SEQ ID NO: 33: COMPUTER: IBM PC Compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/09/253,794 FILING DATE: 2-feb-1999 CLASSIFICATION CUNKNOWN> PRIOR APPLICATION DATA: APPLICATION NUMBER: US 08/318,157 PILING DATE: 05-OCT-1994 ATTORNEY/AGENT INFORMATION: NAME: SAXE, Bernhard D. REGISTRATION NUMBER: 28,665 REFERENCE/DOCKET NUMBER: 18733/464 TELECOMMUNICATION INFORMATION: TELEPHONE: (202)672-5390 TELEPHONE: (202)672-5390 ZIP: 20007-5109 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk CORRESPONDENCE ADDRESS: ADDRESSEE: Foley & Lardner STREET: 3000 K Street, N.W., Suite 500 APPLICANT: HANSEN, Hans J. ARMOUR, Kathryn L. TITLE OF INVENTION: CDR-GRAFTED TYPE III ANTI-CEA HUMANIZED MOUSE MONOCLONAL ANTIBODIES SEQUENCE CHARACTERISTICS: LENGTH: 14 amino acids NUMBER OF SEQUENCES: TYPE: amino acid CITY: Washington STATE: D.C. TELEFAX: (202)672-5399 COUNTRY: USA Score 81; DB 3; : Pred. No. 1.5e-05; 0; Mismatches 0; Length 14;

Indels

0

Gaps

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Sequence 64, Application US/09828708

Patent No. US20020146753A1

GENERAL INFORMATION:
APPLICANT: Ditzel, H.
APPLICANT: Button, D.
APPLICANT: Schaller, M.
APPLICANT: Schaller, M.
TITLE OF INVENTION: Autoantibodies to glucose-6-phosphate isomerase and their particles of INVENTION: Autoimmune disease
FILE REFERENCE: 1361.005US1
CURRENT APPLICATION NUMBER: US/09/828,708
CURRENT FILING DATE: 2001-09-24
NUMBER OF SEQ ID NOS: 123
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 64
LENGTH: 14
TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE 140, Application US/10632706

PUBLICATION NO. US20040175385A1

GENERAL IMPORATION:
APPLICANT: MARKS, JAMES D.
APPLICANT: MARKS, JAMES D.
APPLICANT: MARKS, JAMES D.
TITLE OF INVENTION: THERAPEUTIC MONOCLONAL ANTIBODIES THAT NEUTRALIZE BOTULINUM
TITLE OF INVENTION: NEUROTOXINS
FILE REFERENCE: 407T-895120US
CURRENT FILING DATE: 2003-08-01
CURRENT FILING DATE: 2003-08-01
PRIOR APPLICATION NUMBER: US 60/400,721
PRIOR APPLICATION NUMBER: US 60/400,721
PRIOR APPLICATION NUMBER: US 09/144,806
PRIOR FILING DATE: 2002-08-03
NUMBER OF SEQ ID NOS: 278
SOFTWARE: Patentin version 3.2
RESULT 4
US-10-632-706-144
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US-09-828-708-64
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Best Local S
Matches 14
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                                                                                                                                                                                                                                                                                  OTHER INFORMATION: single chain antibody fragment -10-632-706-140
                                                                                                                                                                                                                                                                                                                             TYPE: PRT
ORGANISM: Artificial
FEATURE:
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                                                                                                                                     1 WVRQAPGKGLEWVA 14
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                                                                                                                                                                                     h 100.0%; Score 81; DB 4; Similarity 100.0%; Pred. No. 1.5e-05; 14; Conservative 0; Mismatches 0;
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TITLE OF INVENTION: THERAPEUTIC MONOCLONAL ANTIBODIES THAT NEUTRALIZE BOTULINUM TITLE OF INVENTION: HEUROTOXINS FILE REFERENCE: 407T-895120US CURRENT APPLICATION NUMBER: US/10/632,706 CURRENT FILING DATE: 2003-08-01 PRIOR APPLICATION NUMBER: US 60/400,721 PRIOR APPLICATION NUMBER: US 60/400,721 PRIOR FILING DATE: 2002-08-01 PRIOR FILING DATE: 2002-08-01 PRIOR APPLICATION NUMBER: US 09/144,806 PRIOR FILING DATE: 1998-08-31 NUMBER OF SEQ ID MOST: 278 SEQ ID MOST: 278 SEQ ID NOS: 278 LENGTH-1.
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RESULT 6
US-10-632-706-152
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US-10-632-706-148
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US-10-632-706-148
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APPLICANT: AAMES D.

APPLICANT: AMERSDORFER, DETER
APPLICANT: AMERSDORFER, PETER
TITLE OF INVENTION: THERAPEUTIC MONOCLONAL ANTIBODIES THAT NEUTRALIZE BOTULINUM
TITLE OF INVENTION: NEUROTOXINS
FILE REFERENCE: 407T-895120US
CURRENT FILING DATE: 105/10/632,706
CURRENT FILING DATE: 2003-08-01
PRIOR APPLICATION NUMBER: US 60/400,721
PRIOR APPLICATION NUMBER: US 60/400,721
PRIOR FILING DATE: 2002-08-01
PRIOR FILING DATE: 1908-08-31
NUMBER OF SEQ ID NOS: 278
SOPTWARE: PAtentin version 3.2
SEQ ID NO 144
                                                                                                                                                                   Query Match
Best Local S
Matches 14
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Publication No. US20040175385A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT
                                                                                                                                                                                                                                                                                             FEATURE:
                                                                                                                                                                                                                                                                                                                ORGANISM: Artificial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 14
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                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%; Score 81; DB 4;
100.0%; Pred. No. 1.5e-05;
                                                                                                                                                                                      100.0%; Score 81; DB 4; 100.0%; Pred. No. 1.5e-05;
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Sequence 152, Application US/10632706 Publication No. US20040175385A1 GENERAL INFORMATION:

TITLE OF INVENTION: THERAPEUTIC MONOCLONAL ANTIBODIES THAT NEUTRALIZE BOTULINUM ITLE OF INVENTION: NEUROTOXINS

MARKS, JAMES D. AMERSDORFER, F

PETER

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; FEATURE:
; OTHER INFORMATION: single chain antibody fragment US-10-632-706-152
                                                                                                                   RESULT 8
US-10-632-706-160
; Sequence 160, Application US/10632706
; Publication No. US20040175385A1
; GENERAL INFORMATION:
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Publication No. US20040175385A1
GENERAL INFORMATION:
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SOFTWARE: PatentIn version 3.2
SEQ ID NO 152
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQ ID NOS: 278
SOFTWARE: PatentIn version 3.2
SEQ ID NO 156
LENGTH: 14
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CURRENT APPLICATION NUMBER: US/10/632,706
CURRENT FILING DATE: 2003-08-01
PRIOR APPLICATION NUMBER: US 60/400,721
PRIOR FILING DATE: 2002-08-01
PRIOR APPLICATION NUMBER: US 09/144,806
PRIOR FILING DATE: 1998-08-31
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CURRENT APPLICATION NUMBER: US/10/632,706
CURRENT FILING DATE: 2003-08-01
PRIOR APPLICATION NUMBER: US 60/400,721
PRIOR FILING DATE: 2002-08-01
PRIOR APPLICATION NUMBER: US 09/144,806
PRIOR FILING DATE: 1998-08-31
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APPLICANT: MARKS, JAMES D.

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APPLICANT: MARKS, JAMES D.
APPLICANT: AMERSDORFER, PETER
TITLE OF INVENTION: THERAPEUTIC MONOCLONAL ANTIBODIES THAT NEUTRALIZE, BOTULINUM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT
ORGANISM: Artificial
FEATURE:
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ORGANISM: Artificial
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Pred. No.
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Pred. No. 1.5e-05;
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CURRENT APPLICATION NUMBER: US/10/632,706
CURRENT FILING DATE: 2003-08-01
PRIOR APPLICATION NUMBER: US 60/400,721
PRIOR EILING DATE: 2002-08-01
PRIOR APPLICATION NUMBER: US 09/144,806
PRIOR APPLICATION NUMBER: US 09/144,806
PRIOR FILING DATE: 1998-08-31
NUMBER OF SEQ ID NOS: 278
SOFTWARE: PATENTIN VETBION 3.2
SEQ ID NO 164
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FILE REFERENCE: 407T-895120US
CURRENT APPLICATION NUMBER: US/10/632,706
CURRENT FILING DATE: 2003-08-01
PRIOR APPLICATION NUMBER: US 60/400,721
PRIOR APPLICATION NUMBER: US 60/400,721
PRIOR FILING DATE: 2002-08-01
PRIOR APPLICATION NUMBER: US 09/144,806
PRIOR FILING DATE: 1998-08-31
NUMBER OF SEQ ID NOS: 278
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                                                                                                                                        Sequence 168, Application US/10632706 Publication No. US20040175385A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Publication No. US20 GENERAL INFORMATION:
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TYPE: PRT
ORGANISM: Artificial
FEATURE:
ENERAL INCURPANCE.

APPLICANT: MARKS, JAMES D.

APPLICANT: AMERSDORFER, PETER

TITLE OF INVENTION: THERAPEUTIC MONOCLONAL ANTIBODIES THAT NEUTRALIZE BOTULINUM

TITLE OF INVENTION: NEUROTOXINS

TITLE OF INVENTION: NEUROTOXINS

7.7.2.2.7.06
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TITLE OF INVENTION: THERAPEUTIC MONOCLONAL ANTIBODIES THAT NEUTRALIZE BOTULINUM
TITLE OF INVENTION: NEUROTOXINS
FILE REFERENCE: 407T-895120US
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OTHER INFORMATION: single chain antibody fragment
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o. US20040175385A1
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100.0%; Pred. No. 1.5e-05;
tive 0; Mismatches 0;
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100.0%; Pred. No. 1.5e-05;
. Mismatches 0;
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Sequence 2, Application US/10489328

| Publication No. US20050037420A1 |
| GENERAL INFORMATION: | GENERATION: | Family device |
| APPLICANT: | Fraunhofer - Gesselschaft zur Forderung der angewandten |
| APPLICANT: | Forschung e.V. |
| APPLICANT: | SCHILLBERG, Stefan |
| APPLICANT: | ZIMMERMANN, Sabine |
| APPLICANT: | ZIMMERMANN, Sabine |
| APPLICANT: | EMAMS, Neil |
| APPLICANT: | EMAMS, Neil |
| APPLICANT: | FISCHER, Rainer |
| TITLE OF INVENTION: | Immunoglobulin having Particular Framework Scaffold and Methods |
| TITLE OF INVENTION of Making and Using |
| FILE REFERENCE: | FRAU-202.1US | | |
| CURRENT | APPLICATION NUMBER: | US/10/409,328 |
| CURRENT | CURRENT | APPLICATION NUMBER: | DCT/US02/29003 |
| PRIOR | APPLICATION NUMBER: | 2004-03-11 |
| PRIOR | APPLICATION NUMBER: | 2004-03-13 |
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Publication No. US20040260068A1
GENERAL INFORMATION:
APPLICANT: TBLUTUSHITA, NAOYA
APPLICANT: Kumar, Shankar
APPLICANT: Vasquez, Maximiliano
TITLE OF INVENTION: Humanied Chicken Antibodies
FILE REFERENCE: 0582.0044.NPUS02
CURRENT APPLICATION NUMBER: US/10/788,625
CURRENT FILING DATE: 2004-02-26
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PRIOR APPLICATION NUMBER: US 60/400,721
PRIOR APPLICATION NUMBER: US 09/144,806
PRIOR APPLICATION NUMBER: US 09/144,806
PRIOR FILING DATE: 1998-08-31
NUMBER OF SEQ ID NOS: 278
SOFTWARE: Patentin version 3.2
SEQ ID NO 168
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US-10-788-625-6
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SOFTWARE: PATENTIN VERSION 3.2
SEQ ID NO 6
LENGTH: 14
TYPE: PRT
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Best Local Similarity
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ORGANISM: Artificial
FEATURE:
OTHER INFORMATION: single chain antibody fragment
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Local Similarity 100.0%; Pred. No. 1.5e-05;
hes 14; Conservative 0; Mismatches 0;
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TITLE OF INVENTION: HUMANIZATION OF ANTIBODIES
FILE REFERENCE: AB600US
CURRENT APPLICATION NUMBER: US/10/923,068
CURRENT FILING DATE: 2004-08-20
NUMBER OF SEQ ID NOS: 518
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 205
LENGTH: 14
TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; ORGANISM: Artificial Sequence ; PEATURE: ; PEATURE: ; OTHER INFORMATION: This is a heavy chain framework region 2 (HFR2) US-10-489-328-2
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US-10-923-068-202
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CURRENT FILING DATE: 2004-08-20
NUMBER OF SEQ ID NOS: 518
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 202
LENGTH: 14
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Publication No. US20050042664A1
GENERAL INFORMATION:
APPLICANT: WL, Herren
APPLICANT: Dall'Acqua, William
APPLICANT: Damschroder, Melissa
TITLE OF INVENTION: HUMANIZATION OF ANTIBODIES
                                                                                                                                                                                                                                                                                                                                        Sequence 205, Applic Publication No. US20 GENERAL INFORMATION:
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SOFTWARE: PatentIn version 3.2
SEQ ID NO 2
Query Match 100.0%;
Best Local Similarity 100.0%;
Matches 14; Conservative 0;
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APPLICANT: Dall'Acqua, William
APPLICANT: Damschroder, Melissa
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Score 81; DB 5;
Pred. No. 1.5e-05;
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Scoring table: Sequence: Title: Perfect score:

13, Appl 15, Appl 16, Appl 16, Appl 16, Appl 49, Appl 50, Appl 17, Appl 18, Appl 25, Appl 25, Appl 25, Appl 26, Appl 26, Appl 26, Appl 26, Appl 26, Appl 27, Appl 62, Appl 63, Appl 64, Appl 65, Appl 66, Appl 67, Appl 68, Appl 69, Appl 60, 
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Total number of hits satisfying chosen parameters:
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1: /cgn2_6/ptodata/2/pubpaa/US10_NEW_PUB.pep:*

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4: /cgn2_6/ptodata/2/pubpaa/US08_NEW_PUB.pep:*

5: /cgn2_6/ptodata/2/pubpaa/US09_NEW_PUB.pep:*

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7: /cgn2_6/ptodata/2/pubpaa/US11_NEW_PUB.pep:*

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Copyright (c) 1993 - 2005 Compugen Ltd.
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FEATURE:
OTHER INFORMATION: Variable heavy chain
OTHER INFORMATION: version 8
US-11-077-978-7
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US-11-077-978-2
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US-11-077-978-7
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APPLICANT: Sherman, Mark A.
APPLICANT: Sherman, Mark A.
APPLICANT: Shively, John B.
APPLICANT: Shively, John B.
APPLICANT: Raubitschek, Andrew A.
APPLICANT: Wu, Anna M.
CURRENT FILING DATE: 2005-03-11
NUMBER OF SEQ ID NOS: 27
SOFTWARB: Patentin version 3.2
SEQ ID NO 7
LENGTH: 120
TYPE: PRT
                                                                                                                                                                                    Sequence 2, Application US/11077978
Publication No. US20050244333A1
GENERAL INFORMATION:
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Publication No. US20050244333A1
GENERAL INFORMATION:
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    APPLICANT: Yazaki, Paul J.
APPLICANT: Sherman, Mark A.
APPLICANT: Shively, John E.
APPLICANT: Raubitschek, Andrew A.
APPLICANT: Wu, Anna M.
TITLE OF INVENTION: A Humanized Anti-CEA T84.66 Antibody and Uses
FILE REFERENCE: 54435.8012.US01
CURRENT APPLICATION NUMBER: US/11/077,978
CURRENT FILING DATE: 2005-03-11
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Sequence 8, Application US/10789273

Publication No. US20050249725A1

GENERAL INFORMATION:
APPLICANT: Basi, Guriq
APPLICANT: Saldanha, Jose
APPLICANT: Yednock, Ted
TITLE OF INVENTION: HUMANIZED ANTIBODIES THAT REC
TITLE OF INVENTION: BETA-AMYLOID PEPTIDE
FILE REFERENCE: ELN-002CP
CURRENT APPLICATION NUMBER: US/10/789,273
CURRENT APPLICATION NUMBER: US/10/386,389
PRIOR APPLICATION NUMBER: US/10/386,389
PRIOR FILING DATE: 2003-03-12
PRIOR APPLICATION NUMBER: US/10/942
PRIOR FILING DATE: 2001-12-06

PRIOR FILING DATE: 2001-12-06

PRIOR FILING DATE: 2001-12-06

PRIOR FILING DATE: 2001-12-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; OTHER INFORMATION: Artificially constructed variable heavy chain ; OTHER INFORMATION: antibody US-11-077-978-3
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US-11-077-978-3
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; OTHER INFORMATION: antibody
US-11-077-978-2
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Best Local S
Matches 14
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APPLICANT: Yazaki, Paul J.
APPLICANT: Shively, John B.
APPLICANT: Shively, John B.
APPLICANT: Raubitschek, Andrew A.
APPLICANT: Raubitschek, Andrew A.
APPLICANT: Wu, Anna M.
APPLICANT: Wu, Anna M.
APPLICANT: Wu, Anna M.
APPLICANTION: A Humanized Anti-CEA T84.66 Antibody and Uses Thereof FILE REFERENCE: 54435.8012.US01
CURRENT APPLICATION NUMBER: US/11/077,978
CURRENT APPLICATION NUMBER: US/11/077,978
CURRENT FILING DATE: 2005-03-11
NUMBER OF SEQ ID NOS: 27
SOFTWARE: Patentin version 3.2
SEQ ID NO 3
LENGTH: 121
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SOFTWARE: PatentIn version 3.2
SEQ ID NO 2
LENGTH: 121
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Best Local :
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ORGANISM: Artificial
FEATURE:
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ORGANISM: Artificial
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APPLICANT: Basi, Guriq
APPLICANT: Saldanha, Jose
APPLICANT: Saldanha, Jose
APPLICANT: Yednock, Ted
TITLE OF INVENTION: HUMANIZED ANTIBODIES THAT RECOGNIZE
TITLE OF INVENTION: BETA-AMYLOID PEPTIDE
FILE REFERENCE: ELM-002CP
CURRENT APPLICATION NUMBER: US/10/789,273
CURRENT FILING DATE: 2004-02-27
PRIOR APPLICATION NUMBER: US/10/388,389
PRIOR FILING DATE: 2003-03-12
PRIOR FILING DATE: 2001-12-06
PRIOR FILING DATE: 2001-12-06
PRIOR FILING DATE: 2000-12-06
PRIOR FILING DATE: 2000-12-06
PRIOR FILING DATE: 2000-12-06
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OTHER INFORMATION: Humanized 3D6 heavy chain variable region FEATURE:
NAME/KEY: SIGNAL
LOCATION: (1)...(19)
US-10-789-273-8
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; LOCATION: (1)...(19)
US-10-789-273-12
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Sequence 27, Application US/10512184

Publication No. US20050244901A1

GENERAL INFORMATION:

APPLICANT: Fraunhofer Gesellschaft zur F"rderung der angewandten Forschung e.V.

TITLE OF INVENTION: Antibodies, recombinant antibodies, recombinant

TITLE OF INVENTION: antibody fragments and fusions mediated plant disease

TITLE OF INVENTION: resistance against fungi
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SOPTWARE: FastSEQ for Windows Version
SEQ ID NO 8
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Best Local Similarity 100.
Matches 14; Conservative
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                                                                                                                                                                                                                                                                                                                                                    Query Match
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SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 13
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ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                 Similarity
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No. US20050249725A1
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ilarity 100.0%;
Conservative (
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Pred. No. 1.4e-06;
); Mismatches 0;
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FILE REFERENCE: 3581.01US01

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CURRENT APPLICATION NUMBER: US/10/512,184
CURRENT FILING DATE: 2004-10-22
NUMBER OF SEQ ID NOS: 72
SOPTWARE: Patentin Ver. 2.1
SEQ ID NO 27
LENGTH: 250
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
         US-10-512-184-72
Sequence 72, Application US/10512184
Sequence 72, Application US/0050244901A1
; GENERAL INFORMATION:
APPLICANT: Fraunhofer Geeellschaft zur F"rderung der angewandten Forschung 'ITILE OF INVENTION: Antibodies, recombinant antibodies, recombinant antibodies, recombinant TITLE OF INVENTION: Antibody fragments and fusions mediated plant disease TITLE OF INVENTION: resistance against fungi
FILE REFERENCE; 3581.01USC;
CURRENT APPLICATION NUMBER: US/10/512,184
; CURRENT PILING DATE: 2004-10-22
; NUMBER OF SEQ ID NOS: 72
; SOFTWARE: Patentin Ver. 2.1
; SEQ ID NO 72
LENGTH: 313
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CURRENT APPLICATION NUMBER: US/10/512,184
CURRENT FILING DATE: 2004-10-22
NUMBER OF SEQ ID NOS: 72
SOPTWARE: PATENTIN VET. 2.1
SEQ ID NO 35
LENGTH: 261
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Publication No. US20050244901A1
GENERAL INFORMATION:
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Best Local Similarity
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Best Local :
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TITLE OF INVENTION: Antibodies, recombinant antibodies, recombinant
TITLE OF INVENTION: antibody fragments and fusions mediated plant disease
TITLE OF INVENTION: resistance against fungi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER INFORMATION: Description of Artificial Sequence: scFv SGB3 OTHER INFORMATION: specificity against Fusarium ssp.; originates OTHER INFORMATION: Gallus gallus.
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TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         100.0%; Score 81; DB 1; 100.0%; Pred. No. 2.3e-06;
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APPLICANT: Basi, Guriq
APPLICANT: Saldanha, Jose
APPLICANT: Saldanha, Jose
APPLICANT: Saldanha, Jose
APPLICANT: Yednock, Ted
ITITLE OF INVENTION: HUMANIZED ANTIBODIES THAT RECOGNIZE
ITITLE OF INVENTION: BETA-AMYLOID PEPTIDE
ITITLE OF INVENTION NUMBER: US/10/789,273
CURRENT APPLICATION NUMBER: US/10/389,389
PRIOR APPLICATION NUMBER: US/10/388,389
PRIOR APPLICATION NUMBER: US/10/388,389
PRIOR FILING DATE: 2001-12-06
PRIOR APPLICATION NUMBER: US 10/010,942
PRIOR APPLICATION NUMBER: US 60/251,892
PRIOR PILING DATE: 2000-12-06
NUMBER OF SEQ ID NOS: 63
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 10
LENGTH: 98
TWORD: DOT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER INFORMATION: Description of Artificial Sequence: scFv FPCWPA5; OTHER INFORMATION: with specificity against Pusarium ssp.; originat; OTHER INFORMATION: from Gallus gallus.
US-10-512-184-28
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LENGTH: 252
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Publication No. US20050244901A1
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FILE REFERENCE: 3581.01US01
CURRENT APPLICATION NUMBER: US/10/512,184
CURRENT FILING DATE: 2004-10-22
NUMBER OF SEQ ID NOS: 72
NUMBER OF SEQ ID NOS: 72
NUMBER OF SEQ ID NOS: 72
NUMBER: Patentin Ver. 2.1
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Best Local Similarity
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Similarity 100.0%; Pred. No. 2.7
14; Conservative 0; Mismatches
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10. US20050249725A1
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Pred. No.
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; ORGANISM: Homo sapiens
US-10-789-273-10
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                                                                                                                         FILE REFERENCE: ELN-002CP
CURRENT APPLICATION NUMBER: US/10/789,273
CURRENT FILING DATE: 2004-02-27
PRIOR APPLICATION NUMBER: US/10/388,389
PRIOR FILING DATE: 2003-03-12
PRIOR APPLICATION NUMBER: US 10/010,942
PRIOR FILING DATE: 2001-12-06
PRIOR APPLICATION NUMBER: US 60/251,892
PRIOR FILING DATE: 2000-12-06
                                                                                                                                                                                                                                                                                                                     Sequence 9, Application US/10789273

Publication No. US20050249725A1

GENERAL INFORMATION:
APPLICANT: Basi, Guriq
APPLICANT: Saldanha, Jose
APPLICANT: Yednock, Ted
APPLICANT: Yednock, Ted
TITLE OF INVENTION: HUMANIZED ANTIBODIES THAT RECOGNIZE
TITLE OF INVENTION: BETA-AMYLOID PEPTIDE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 92.9
Matches 13; Conservative
                      NUMBER OF SEQ ID NOS: 63
SOPTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 9
LENGTH: 121
TYPE: PRT
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 32, Application US/11144248 Publication No. US20050244408A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: US/10/038,591
PRIOR PILING DATE: 2002-01-04
PRIOR APPLICATION NUMBER: 60/259,927
PRIOR PILING DATE: 2001-01-05
NUMBER OF SEQ ID NOS: 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/11/144,248
CURRENT FILING DATE: 2005-06-02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Miller, Penelope B.
APPLICANT: Moyer, James D.
APPLICANT: Corvalan, Jose R.
APPLICANT: Garlo, Michael
TITLE OF INVENTION: ANTIBODIES TO INSULIN-LIKE GROWTH FACTOR I RECEPTOR FILS REFERENCE: ABX-PF2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: PatentIn Ver.
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APPLICANT: Beebe, Jean
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        y Match 96.3%;
Local Similarity 92.9%;
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92.9%;
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Pred. No. 2.
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JS-11-144-248-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 8, Application US/11144248 Publication No. US20050244408A1 GENERAL INFORMATION:
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                                                                                                SEQ ID NO 8
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Best Local :
                                                                                                                                                           APPLICANT: Miller, Penelope E.
APPLICANT: Moyer, James D.
APPLICANT: Moyer, James D.
APPLICANT: Corvelan, Jose R.
APPLICANT: Gallo, Michael
TITLE OF INVENTION: ANTIBODIES TO INSULIN-LIKE GROWTH FACTOR I RECEPTOR FILE REFERENCE: ABX-PF2
CURRENT APPLICATION NUMBER: US/11/144,248
CURRENT FILING DATE: 2005-06-02
PRIOR APPLICATION NUMBER: US/10/038,591
PRIOR APPLICATION NUMBER: US/10/038,591
PRIOR APPLICATION NUMBER: US/10/038,591
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PRIOR APPLICATION NUMBER: US/10/038,591
PRIOR APPLICATION NUMBER: US/10/038,591
PRIOR APPLICATION NUMBER: US/10/038,591
PRIOR APPLICATION NUMBER: 00/259,927
PRIOR APPLICATION NUMBER: 00/259,927
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Cohen, Bruce D.
APPLICANT: Beebe, Jean
APPLICANT: Miller, Penelope E.
APPLICANT: Moyer, James D.
APPLICANT: Gallo, Michael
APPLICANT: Gallo, Michael
TITLE OF INVENTION: ANTIBODIES TO INSULIN-LIKE GROWTH FACTOR I RECEPTOR FILE REFERENCE: ABX-PF2
CURRENT APPLICATION NUMBER: US/11/144,248
CURRENT FILING DATE: 2005-06-02
PRIOR APPLICATION NUMBER: US/10/038,591
PRIOR PILING DATE: 2002-01-04
PRIOR APPLICATION NUMBER: 60/259,927
REFOR EFILING DATE: 2001-05
                                                                                                                   PRIOR FILING DATE: 2001-01-05
NUMBER OF SEQ ID NOS: 60
SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: Patentin Ver. 2.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Cohen, Bruce D. APPLICANT: Beebe, Jean
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Homo sapiens
                        ORGANISM: Homo sapiens
                                              LENGTH: 124
TYPE: PRT
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Local Similarity 92.9%;
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Pred. No.
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Pred. No. 3.4e-06;
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Oberty Match

Best Local Similarity

95.3%, Score 78; DB 7; Length 124;

Best Local Similarity

92.9%, Pred. Mo. 3.4e-06;

Macches 13) Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Macches 13) Conservative 48

RESULT 15

US-11-144-248-16

[WS-11-144-248-16

[Sequence 15, Application US/11144248

PREDIAL MINOSWATION

APPLICANT, Cohen, Bauce D.

APPLICANT, Miller, Penelope E.

APPLICANT, Moyer, James D.

APPLICANT, MALLER, Penelope E.

APPLICANT, Moyer, James D.

APPLICANT, MOYER, James D.

APPLICANT, MALLER, Penelope E.

APPLICANT, MOYER, US/11/144, 248

CURRENT APPLICATION MOMERS, US/11/144, 248

CURRENT APPLICATION MOMERS, US/10/038, 591

PRIOR APPLICATION MOMERS, US/10/038, 591

PRIOR FILING DATE, 2001-01-05

INDELS OF SEQ ID MOX: 60

I SEQ ID MOX 16

I PRIOR FILING DATE, 2001-01-05

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Result
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Maximum DB
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Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.
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seq length: 2000000000
protein search, using sw model
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Copyright (c) 1993 - 2005 Compugen Ltd.
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## ALIGNMENTS

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RESULT 1
ADR38751
antibacterial; antibody; botulinum neurotoxin type A; BoNT/A; BoNT/A neutralising epitope; anti-botulinum neurotoxin antibody; toxin neutralisation; botulinum neurotoxin poisoning; mouse; heavy chain; CDR2; complementarity determinig region 2.
                                                                                                                                                                           31-AUG-1998; 98US-00144886.
01-AUG-2002; 2002US-0400721P.
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                                                                                                                                                            (REGC ) UNIV CALIFORNIA.
                                                                                                                                                                                                                   09-SEP-2004.
                                                                                                                                                                                                                                                                                                         chain anti-BoNT-antibody CD2
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The invention describes an isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a specific clone where (I) binds to and neutralises botulinum neurotoxin type A (BoNT/A). An isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a clone chosen from clone 25, C35, C35, 1C6, 3D12, B4, 1F3, huC25, A1, Ar2, WR1(V), WR1(T), 3-1, 3-8, 3-10 and ING1, where (I) binds to and neutralizes botulinum neurotoxin type A (BoNT/A). Also described are: a polypeptide (II) comprising BoNT/A neutralising epitope having an epitope that is

New isolated antibody that neutralizes botulinum neurotoxin type A, useful for diagnosing botulism or for treating pathologies associated with botulinum neurotoxin poisoning.

Example 4; SEQ ID NO 153; 110pp; English

WPI; 2004-652009/63.

Marks JD,

Amersdorfer

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Matches 17
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The invention describes an isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a specific clone where (I) binds to and neutralises botulinum neurotoxin type A (BONT/A). An isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a clone chosen from clone S25, C25, C39, 1C6, 3D12, B4, 1F3, huC25, Ar1, Ar2, WR1(Y), WR1(T), 3-1, 3-8, 3-10 and ING1, where (I) binds to and neutralizes botulinum

The invention describes an isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a specific clone where (I) binds to and neutralises botulinum neurotoxin type A (BoNT/A). An isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a clone chosen from clone

Example

4

SEQ ID NO 141; 110pp; English

Example 4;

SEQ ID NO 145; 110pp; English.

botulinum neurotoxin poisoning.

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Matches 17
                                                                                                                                                                                                                                                                 New isolated antibody that neutralizes bbtulinum neurotoxin type A, useful for diagnosing botulism or for treating pathologies associated with botulinum neurotoxin poisoning.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Marks
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01-AUG-2002; 2002US-0400721P.
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Pred. No.
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01-AUG-2002; 2002US-0400721P.
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                                                                                                                                                                                                                                                           New isolated antibody that neutralizes botulinum neurotoxin type A, useful for diagnosing botulism or for treating pathologies associated
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Marks JD,
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                                                                                                                                                                                                                               botulinum neurotoxin poisoning.
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Pred. No. 1.2e-07;
Mismatches 0; Indels
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The invention describes an isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a specific clone where (I) binds to and neutralises botulinum neurotoxin type A.

The invention describes an isolated antibody (I) that specifically binds

Example 4; SEQ ID NO 157; 110pp; English

Example 4; SEQ ID NO 149; 110pp; English

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RESULT 5
ADR38755
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                                                                 New isolated antibody that neutralizes botulinum neurotoxin type A, useful for diagnosing botulism or for treating pathologies associated with botulinum neurotoxin poisoning.
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01-AUG-2002; 2002US-0400721P.
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100.0%; Pred. No.
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RESULT 6
ABU56860
ID ABU5
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Best Local Similarity
Matches 17; Conserv
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                                 Novel antibody that specifically binds and neutralizes botulinum neurotoxin type A useful for neutralizing botulinum neurotoxin and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ds snw
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Botulinum neurotoxin type A; BoNT/A; mouse; heavy chain variable region; scFv; antibody; botulism; antibacterial; single chain antibody; VH;
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treating botulism.
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                                                                                                                                                                                                                                                                                              (MARK/) MARKS J D.
(AMER/) AMERSDORFER P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          immunoglobulin.
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                                                                                                                                                                                                                          Amersdorfer
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밁 8

Claim

6; Page 22; 31pp; English

The invention relates to an isolated antibody that specifically binds to an epitope specifically bound by an antibody expressed by a clone such as

Claim 6; Page 22; 31pp; English

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RESULT 7
ABU56861
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local S
Matches 17
                                                                                                     Novel antibody that specifically binds and neutralizes botulinum neurotoxin type A useful for neutralizing botulinum neurotoxin autreating botulism.
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                                                                                                                                                                                                                                                                                                                                                                                                 31-AUG-1998;
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                                                                                                                                                                                               WPI; 2003-182618/18.
                                                                                                                                                                                                                                         Marks JD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         immunoglobulin.
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(AMER/) AMERSDORFER P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         h 100.0%; Score 93; DB 6; Similarity 100.0%; Pred. No. 5.7e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 binding antibody scTv VH region from C25
                                                                                                                                                                                                                                           Amersdorfer P;
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                                                                                                                                                                                                                                                                                                                                                                                                   98US-00144886.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mouse heavy chain variable region scrv seqid 59.
                                                                                                                                                                                                                                         New isolated antibody that neutralizes botulinum neurotoxin type A, useful for diagnosing botulism or for treating pathologies associated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              31-AUG-1998; 98US-00144886.
01-AUG-2002; 2002US-0400721P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-AUG-2003; 2003US-00632706.
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                                                                                                                                                                                                                                                                                                                                            WPI; 2004-652009/63.
                                                                                                                                                                                                                                                                                                                                                                                                            Marks JD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    heavy chain variable region; single chain antibody; scFv.
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                                                                                                                                                                                                         botulinum
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l Similarity 100.0%;
17; Conservative (
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                                                                                                                                                                                                      neurotoxin poisoning.
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The invention describes an isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a specific clone where (I) binds to and neutralises botulinum neurotoxin type A

Example 1;

SEQ ID NO 59; 110pp; English.

New isolated antibody that neutralizes botulinum neurotoxin type A, useful for diagnosing botulism or for treating pathologies associated with botulinum neurotoxin poisoning.

WPI; 2004-652009/63

Marks JD,

Amersdorfer P;

(REGC ) UNIV CALIFORNIA.

31-AUG-1998; 01-AUG-2002;

98US-00144886. 2002US-0400721P.

01-AUG-2003; 2003US-00632706.

Example 1; SEQ ID NO

60; 110pp; English

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RESULT 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CC (BoNT/A). An isolated antibody (I) that specifically binds to an epitope CC specifically bound by an antibody expressed by a clone chosen from clone CS 25, C25, C39, IC6, 3D12, B4, IF3, huC25, Ar1, Ar2, WR1(Y), WR1(T), 3-1, CC comprising BONT/A neutralising epitope dare: a polypeptide (II) CC specifically bound by an antibody expressed by clones as mentioned in (I) producing (I); and a composition (III) comprising several anticc different epitope of a botulinum neurotoxin antibodies, where each antibody is specific for a CC different epitope of a botulinum neurotoxin and the combination of CC comprising (I); and a kit comprising (I) is useful for neutralising a comprising (I); and a kit comprising (I) is useful for neutralising a botulinum neurotoxin which CC comprising (I); and a kit comprising a botulinum neurotoxin which CC involves contacting neurotoxin with (I) in surplus, where each of (I) is specific for a different epitope of the botulinum neurotoxin and the combination of antibodies shows greater toxin neutralisation than the single antibodies in surplus. (I) in surplus, where each of (I) is specific for a different epitope of the botulinum neurotoxin and the combination of antibodies shows greater toxin neutralisation than the single antibodies in surplus. (I) is useful for diagnosing the botulism or for treating pathologies associated with botulinum neurotoxin composition or diagnosis of botulism. This is the amino acid equence of a mouse heavy chain variable fragment anti-botulinum toxin
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                                                                                                                                                                                                                                                                                                                                                                                                               antibacterial; antibody; botulinum neurotoxin type A; BONT/A; BONT/A neutralising epitope; anti-botulinum neurotoxin antibody; toxin neutralisation; botulinum neurotoxin poisoning; mouse; heavy chain variable region; single chain antibody; scFv.
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         WPI; 1994-167486/20
                                                   Tempest PR,
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Sequences are provided for mouse 022 VH (AAR54931), humanized NEWM-based VH (022 KUVH, AAR54929), humanized KOL-based VH (022 KUVH, AAR54930), mouse 022 VK (AAR54933) and humanized REI-based VK (022 HUVK, AAR54933)

During hab production, VH and VK cDNAs were PCR amplified using primers given in AAQ65378-87. Mutagenesis of clone M13VHPCR2 KOL VH (L71R) was performed using oligos AAQ65388-89. The habs can be used in heteroantibody, bifunctional antibody and immunocoxin production. (Updated on 25-MAR-2003 to correct PN field.) (Updated on 27-AUG-2003 to
                                                                                                                                                                                                                                                                                                                                                                                  Fc receptor; FCR; humanized antibody; hAb; IgG; cancer; allergy; autoimmune disease; heteroantibody; bifunctional antibody; immunotoxin; CDR; complementarity determining region; WH; heavy chain variable regio VK; kappa chain variable region; mononuclear phagocyte; PCR;
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            Disclosure; Page 14;
                                         diseases.
                                                                     New humanised antibodies
                                                                                                  WPI; 1994-167486/20.
                                                                                                                              Tempest PR,
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Sequences are provided for mouse 022 VH (AAR54931), humanized NEWM-based VH (022 KUVH, AAR54929), humanized KOL- based VH (022 KUVH, AAR54930), mouse 022 VK (AAR54933) and humanized REI- based VK (022 KIVH, AAR54930).

During hab production, VH and VK CDNAs were PCR amplified using primers given in AAQ6538-97. Mutagenesis of clone M13VHPCR2 KOL VH (L71R) was performed using oligos AAQ65388-89. The hAbs can be used in heteroantibody, bifunctional antibody and immunotoxin production. (Updated on 25-MAR-2003 to correct PN field.) (Updated on 27-AUG-2003 to
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Humanized antibodies (hAbs) for IgG Fc receptors on human phagocytes comprise the CDR of mouse monoclonal antibody 22 (from hybridoma 022WCL-1), VH chains from human Igs NEWM or KOL, and VK chains from Ig REI. Sequences are provided for mouse 022 VH (AAR5491), humanized NDL-based VH (022 NMVH, AAR54929), humanized KOL-based VH (022 KLVH, AAR54930), mouse 022 VK (AAR54933) and humanized REI-based VK (022 KUVK, AAR54930). During hab production, VH and VK CDNAs were PCR amplified using primers given in AA65378-87. Mutagenesis of clone M13VHPCR2 KOL VH (L71R) was performed using oligos AAQ65388-89. The hAbs can be used in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Fc receptor; FcR; humanized antibody; hAb; IgG; cancer; allergy; autoimmune disease; heteroantibody; bifunctional antibody; immunotoxin;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  antibody; MAb.
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(first_entry)
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Pred. No. 1.1e-06;
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RESULT 14
AAO18397
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AC AAO18

AAO18397 standard; protein; 120 AA.

AA018397;

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ARO RAD18
ARC ARO18
ARC AR
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                                                                                                                                                                                                                                                      The present invention relates to humanised anti-lymphotoxin beta receptor (LT-beta-R) antibodies. These are derived from the murine LT-beta-R binding antibody CBE11 and can be used to treat neoplasia in humans. The present sequence is the murine CBE11 heavy chain variable region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New humanized anti-lymphotoxin-beta receptor antibody, useful for treating or reducing the advancement, severity or effects of neoplasia, particularly solid tumors (i.e. carcinomas) including colorectal cancer
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13-MAR-2001; 2001US-0275289P
21-JUN-2001; 2001US-0299987P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mouse; humanised antibody; lymphotoxin beta receptor; antibody; cancer; neoplasia; LT-beta-R; light chain; heavy chain; variable region.
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                                                                                                                                                                                                        Sequence 120 AA;
                                                                                                                                                                                                                                                                                                                                                                                        Claim 2; Page 9-10; 41pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2002-583337/62.
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17; Conserv
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                                                   TISDGGSYTYYPDSVKG 17
     TISDGGSYTYYPDSVKG
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Pred.
                                                                                                                               Score 93; DB 5;
Pred. No. 1.1e-06;
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RESULT 15
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CX Huma
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New humanized anti-lymphotoxin-beta receptor antibody, useful for treating or reducing the advancement, severity or effects of neoplasia, particularly solid tumors (i.e. carcinomas) including colorectal cancer
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13-MAR-2001; 2001US-0275289P.
21-JUN-2001; 2001US-0299987P.
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                                                                                                                                                                                                                                                                                    Mouse; humanised antibody; lymphotoxin beta receptor; antibody; cancer; neoplasia; LT-beta-R; light chain; heavy chain; variable region.
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N-PSDB; AAL48454.
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                                          12-OCT-2001; 2001WO-US032140
                                                                                             18-APR-2002.
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                                                                                                                                                                                                     Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                   murine CBE11 heavy chain variable domain #1.
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Matches 17
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                                                                                                                                                                                                                                                                                                     New humanized anti-lymphotoxin-beta receptor antibody, useful for treating or reducing the advancement, severity or effects of neoplasia, particularly solid tumors (i.e. carcinomas) including colorectal cancer and breast cancer.
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13-MAR-2001; 2001US-0275289P.
21-JUN-2001; 2001US-0299987P.
                                                                                                                                                                        Sequence 120 AA;
                                                                                                                                                                                                                                                                         Disclosure; Page 14; 41pp; English.
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N-PSDB; AAL48452.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 (BIOJ ) BIOGEN INC.
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                                                       1 TISDGGSYTYYPDSVKG
                                                                                                                             h 100.0%; Score 93; DB 5;
Similarity 100.0%; Pred. No. 1.1e-06;
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and is derived by analysis of the total score distribution.
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Copyright (c) 1993 - 2005 Compugen Ltd.
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A;Molecule type: mRNA
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A;Residues: 1-98 <LEV>
A;Cross-references: UNIPROT:P18528; UNIPARC:UPI00000278DB
A;Cross-references: UNIPROT:P18528; UNIPARC:UPI00000278DB
A;Experimental source: strain BALB/cJ
A;Note: this sequence belongs to the VH7183 subfamily
C;Superfamily: immunoglobulin V region; immunoglobulin
C;Keywords: heterotetramer; immunoglobulin
P;15-98/Domain: immunoglobulin homology <IMM>
P;15-98/Domain: immunoglobulin homology <IMM>
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71.0	71.0	72.0	72.0	72.0	74.2	74.2	74.2	75.3	75.3	77.4	78.5	78.5	78.5	79.6	79.6
117	113	119	118	117	124	118	111	123	108	112	121	117	114	138	118
μ	N	N	N	μ	N	N	N	N	N	N	N	μ	N	N	N
HVMS34	S26468	A43413	S00700	HVMS84	I27888	S20641	PH1007	G27888	PH1006	A27889	H27888	HVMS39	PH1009	S09258	PH0097
Ig heavy	Ig heavy	Ig heavy				_	_	_		_					Ig heavy
y chain	y chain	y chain	y chain	y chain	_	_	-	-		y chain	y chain	y chain pre	chain	y chain	chain

## ALIGNMENTS

R;Levy, N.S.; Malipiero, U.V.; Lebecque, S.G.; Gearhart, P.J. Exp. Med. 169, 2007-2019, 1989
A;TItle: Early onset of somatic mutation in immunoglobulin VH A;Reference number: JT0501; MUID:89279149; PMID:2499654
A;Accession: JT0501

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Length 98

Ig heavy chain V region (6.96) - mouse C;Species: Mus musculus (house mouse) C;Date: 30-Jun-1990 #sequence\_revision 30-Jun-1990 #text\_change 09-Jul-2004 C;Accession: JT0501

HVMS96

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A;Cross references: UNIPARC:UPI0000176A0C
A;Experimental source: strain Balb/c
A;Experimental source: strain Balb/c
A;Experimental source: strain Balb/c
A;Note: this sequence was determined from the germline gene
C;Comment: This chain was isolated from a hybridoma protein that binds influenza virus he
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
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A;Title: Structural and functional implications of a restricted antibody response A;Reference number: A91043; MUID:86300658; PMID:2427335
A;Accession: C27889
                                                                                                                                                                                                                                                                                                                                                         Ig heavy chain V region (H220-25) - mouse C;Species: Mus musculus (house mouse) C;Species: Mus musculus (house mouse) C;Date: 15-Dec1-1988 #sequence_revision 15-Ded-1988 #text_change 16-Aug-1996 C;Accession: C27889
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A; Residues: 1-101 < CAT>
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  Score 93;
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     Length 101;
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R;Lebecque, S.G; Gearhart, P.J.

J. Exp. Med. 172, 1717-1727, 1990
A;Title: Boundaries of somatic mutation in rearranged immunoglobulin genes: 5' boun A; Reference number: $70410; MUID:91079775; PMID:2258702
A;Accession: $70410
A;Accession: $70410
A;Status: translation not shown
A;Molecule type: DNA
A;Residues: 1-19 <LEB>
A;Cross-references: UNIPARC:UPI0000115881; EMBL:X53776; NID:g52475; PIDN:CAA37792.1
C;Genetics:
A;Introns: 16/1
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;1-19/Domain: signal sequence #status predicted <SIG>
F;20-152/Product: Ig heavy chain V region MAK33 #status predicted <MAT>
F:34-117/Tomain: ifmonoriobulin homology
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A;Cross-references: UNIPARC:UPI0000176B67
A;Cross-references: UNIPARC:UPI0000176B67
A;Bxperimental source: strain Balb/c
A;Note: this sequence was determined from the germline gene
C;Comment: This chain was isolated from a hybridoma protein that binds influenza virus
C;Superfamily: immunoglobulin vregion; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;15-98/Domain: immunoglobulin homology <IMM>
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A;Title: Structural and functional implications of a restricted A;Reference number: A91043; MUID:86300658; PMID:2427335
A;Accession: B27889
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A;Title: Cloning and nucleotide sequence of heavy- and A;Reference number: A91572; MUID:87248058; PMID:3110009
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ig heavy chain precursor V region (MAK33) - mouse C;Species: Mus musculus (house mouse) C;Date: 05-Jun-1988 #sequence revision 05-Jun-1988 C;Accession: B26471; S70410
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                                                                                Length 152;
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A; Molecule type: mRNA
A; Residues: 1-102 < CH
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Richen, Q.; Stenzel-Poore, M.; Rittenberg, M.B. submitted to the EMBL Data Library, March 1991 A;Description: Natural polyreactive antibodies (A;Reference number: $14484 A;Accession: $14580 A;Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       R;Lawler, A.M.; Lin, P.S.; Gearhart, P.J.
Proc. Natl. Acad. Sci. U.S.A. 84, 2454-2458, 1987
A;Title: Adult B-cell repertoire is biased toward two heavy-chain A;Reference number: A94148; MUID:87175692; PMID:3104915
A;Accession: C25913
A;Mcolecule type: DNA
A;Residues: 1-83 <LAW>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              A,Cross-references: UNIPARC:UPI0000115F39; EMBL:X58652; NID:g51293; PIDN:CAA41509.1; C;Superfamily: immunoglobulin V region; immunoglobulin homology C;Keywords: heterotetramer; immunoglobulin F;7-90/Domain: immunoglobulin homology <IMM>
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C;Species: Mus musculus (house mouse)
C;Date: 20-Peb-1995 #sequence_revision
C;Accession: S14580
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C;Superfamily: immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
                                                               C;Accession: S14581
R;Chen, Q.; Stenzel-Poore, M.; Rittenberg, M.B. submitted to the RMEL Data Library, March 1991
A;Description: Natural polyreactive antibodies
                                                                                                                                             Ig heavy chain V region - mouse
C;Species: Mus musculus (house mouse)
C;Date: 20-Feb-1995 #sequence_revision
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A; Residues: 1-94 < CHE>
A, Status: preliminary
                       A; Reference number: A; Accession: S14581
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94.1%; Pred. No.
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A;Cross-references: UNIPARC:UPI0000176B68
A;Experimental source: strain Balb/c
A;Note: this sequence was determined from the germline gene
C;Comment: This chain was isolated from a hybridoma protein that binds influenza virus
C;Superfamily: immunoglobulin v region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
C;Keywords: heterotetramer; immunoglobulin
F;15-98/Domain: immunoglobulin homology
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J. Exp. Med. 176, 761-779, 1992
A;Title: Both IgM and IgG anti-DNA antibodies are the products of clonally selective A;Reference number: PH0971; MUID:92381444; PMID:1512540
A;Accession: PH1010
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R;Caton, A.J.; Brownlee, G.G.; Staudt, L.M.; Gerhard, W. EMBO J. 5, 1577-1587, 1986
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A;Cross-references: Unicono176D1A
A;Cross-references: B cell, strain [NZB x NZW]F1
A;Experimental source: B cell, strain [NZB x NZW]F1
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
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A; Residues: 1-119 < CAT>
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C;Species: Mus musculus (house mouse)
C;Date: 15-Dec-1988 #sequence_revision 15-Dec-1988 #text_change 16-Aug-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Status: nucleic acid sequence not shown molecule type: mRNA
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;Species: Mus musculus (house mouse)
;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change
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94.1%;
                                                                                                    93.5%; Score 87; DB 2; LE
94.1%; Pred. No. 3.8e-06;
94.1%; Omismatches 1;
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Pred. No. 3.2e-06;
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Pred. No. 3.4e-0
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                        A; Molecule type: DNA
A; Residues: 1-122 < CAT>
A;Cross-references: UNIPARC:UPI0000176B6B
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A;Residues: 1-120 <BOB>
A;Residues: 1-120 <BOB>
A;Cross-references: UNIPARC:UPI0000116200; EMBL:X82589; NID:g854304; PIDN:CAA57925.1; PII
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
C;Keywords: heterotetramer; immunoglobulin
F;14-97/Domain: immunoglobulin homology <IMM>
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C;Species: Mus musculus (house mouse)
C;Date: 27-Oct-1995 #sequence_revision 03-Nov-1995 #text_change 23-Jul-1999
C;Accession: S55537
                                                                                                                            Ig heavy chain V region (H35-C6) - mouse
C;Species: Mus musculus (house mouse)
C;Date: 15-Dec-1988 #sequence_revision 15-Ded-1988 #text_change 16-Aug-1996
C;Accession: E27888
                                                                                                                                                                                                                                             RESULT 12
E27888
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A;Reference number: S55528; MUID:95239763; PMID:7536850
A;Accession: S55536
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ig heavy chain V region pe20 - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 27-Oct-1995 #sequence_revision 03-Nov-1995 #text_change 23-Jul-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Cross-references: UNIPARC:UDI0000116201; EMBL:X82590; NID:g854306; PIDN:CAA57926.1; PII C;Superfamily: immunoglobulin V region; immunoglobulin homology C;Keywords: heterotetramer; immunoglobulin F;14-97/Domain: immunoglobulin homology <IMM>
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A; Residues: 1-120 < BOE>
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J. Mol. Biol. 247, 932-946, 1995
A;Tille: Comprehensive epitope analysis of monoclonal anti-proenkephalin antibodies using utations in the variable region genes.
R;Caton, A.J.; Brownlee, G.G.; Staudt, L.M.; Gerhard, W.
EMBO J. 5, 1577-1587, 1986
A;Title: Structural and functional implications of a restricted antibody response to A;Reference number: A91043; MUID:86300658; PMID:2427335
A;Accession: E27888
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R;Boettger, V.; Boettger, A.; Lane, E.B.; Spruce, B.A
J. Mol. Biol. 247, 932-946, 1995
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                                                                                                        G.G.; Staudt, L.M.; Gerhard, W.
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Pred. No. 3.8e-06;
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Pred. No. 3.8e-06;
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Ricaton, A.J.; Brownlee, G.G.; Staudt, L.M.; Gerhard, W.
EMBO J. 5, 1577-1587, 1986

A,Title: Structural and functional implications of a restricted antibody response to a (A,Title: Structural and functional implications of a restricted antibody response to a (A,Title: Structural and functional implications of a restricted antibody response to a (A,Title: Structural and functional implications of a restricted antibody response to a (A,Title: Structural and functional implications of a restricted antibody response to a (A,Title: Structural)
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A;Res
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S63597
Ig heavy chain, V region - mouse
C;Species: Mus musculus (house mouse)
C;Date: 28-Oct-1996 #sequence_revision 13-Mar-1997 #text_change 21-Jan-2000
C;Date: 28-Oct-1996 #sequence_revision 13-Mar-1997 #text_change 21-Jan-2000
C;Accession: S63597
R;Verdaguer, N.; Mateu, M.G.; Bravo, J.; Domingo, E.; Fita, I.
J. Mol. Biol. 256, 364-376, 1996
A;Title: Induced pocket to accommodate the cell attachment Arg-Gly-Asp motif: A;Reference number: S63596; MUID:96174482; PMID:8594203
A;Accession: S63597
A;Status: preliminary; not compared with conceptual translation
A;Molecule type: mRNA
A;Molecule type: mRNA
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/Species: Mus musculus (house mouse)
/Date: 15-Dec-1988 #sequence_revision 15-Dec-1988 #text_change 16-Aug-1996
/Accession: 127887
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Similarity 94.1%;
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Pred. No. 3.9e-06;
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Pred. No. 1.1e-05;
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RESULT 15

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Ig heavy chain V region (H37-62) - mouse (FSpecies: Mus musculus (house mouse) (FSpecies: Mus musculus (house mouse) (FSpecies: Mus musculus (house mouse) (FSpecies: 15-Dec-1988 #sequence_revision 15-Dec-1988 #text_change 16-Aug-1996 (FAccession: C27888 R.Caton, A.J.; Brownlee, G.G.; Staudt, L.M.; Gerhard, W. R.Caton, A.J.; Brownlee, G.G.; Staudt, L.M.; Gerhard, W. R.Fitle: Structural and functional implications of a restricted antibody response to a de A.Fitle: Structural and functional implications of a restricted antibody response to a A.Feference number: A91043; MUID:88300658; PMID:2427335
                                                                                                                                                                                                                                                   A;Molecule type: DNA
A;Residues: 1-124 <CAT>
A;CROSE-references: UNIPARC:UPI0000176B71
A;Experimental source: strain Balb/c
A;Note: this sequence was determined from the germline gene
C;Comment: This chain was isolated from a hybridoma protein that binds influenza virus he
C;Superfamily: immunoglobulin v region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;15-98/Domain: immunoglobulin homology <IMM>
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                                                                                                                         Query Match
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Matches 15; Conserva
   50 TISSEGTYTYYPDSVKG
                                     1 TISDGGSYTYYPDSVKG 17
                                                                                                                         Conservative
                                                                                                                                                           90.3%; Score 84; DB 2;
88.2%; Pred. No. 1.1e-05;
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Search completed: November 21, 2005, 12:22:15 Job time: 4.14103 secs

OM protein - protein search, using sw

mode1

GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.

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November 21, 2005, 11:45:55; Search time 25.1575 Seconds (without alignments)
476.756 Million cell updates/sec

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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Q65ZL8 MOUSE
HV58 MOUSE
HV59 MOUSE
HV59 MOUSE
HV59 MOUSE
HV55 MOUSE
Q5XHD5 XENLA
HV55 MOUSE
Q4T136 TETNG
Q5EK12 FRAT
Q4VEH1 RAT
Q6EZIG HUMAN
Q4T5M5 TETNG
Q5EX12 FRAT
Q4T5M5 TETNG
Q5EX12 FRAT
Q4T5M5 TETNG
Q5EX12 KENLA
HV3L HUMAN
Q5FZTB MOUSE
Q6EDFOT XENLA
HV3L HUMAN
Q5FZTB MOUSE
Q6EDFOT MOUSE
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Q65218 mus musculu
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P18525 mus musculu
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Q5xhd5 xenopus lae
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Q5pk9 rattus norv
Q4f584 retraodon n
Q5pk9 rattus norv
Q4f584 xenopus lae
P18524 mus musculu
Q6ddq7 xenopus lae
P01773 homo sapien
Q5f218 mus musculu
Q5m8x4 xenopus tro
Q91207 mus musculu
Q91171 homo sapien
Q5f3f4 homo sapien
Q5f3f4 homo sapien
Q5f3f6 rattus norv
Q96bb9 homo sapien
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Description
 RESULT 2
Q99KA4 MOUSE
D Q99KA4_MOUSE PRELIMINARY;
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31 31 31 31 31 31 31 31 31 31 31 31 31

PRT;

487 AA.

Regult No.

Score

Query Match

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database

2:

Minimum DB Maximum DB

seq length: 0
seq length: 2000000000

Total number of hits satisfying chosen parameters:

2166443

2166443 segs, 705528306 residues

Scoring table: Sequence: Perfect score:

BLOSUM62 Gapop 10.0 , Gapext 0.5

US-10-632-706-157 93

1 TISDGGSYTYYPDSVKG

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1 TISDGGSYTYYPDSVKG 17                50 TISDGGSYTYYPDSVKG 66	ry Match 100.0%; Score 93; DB 1; t Local Similarity 100.0%; Pred. No. 1.4e-06; ches 17; Conservative 0; MismatcHes 0;	PIR; JT0501; HYWS96.  HSSP; P01783; 11GC.  SMR; P18528; 1-98.  InterPro; IPR007110; Ig-like.  InterPro; IPR0073596; Ig_v.  SMART; SM00406; IGv; 1.  PROSITE; PS50835; IG_LIKE; 1.  Immunoglobulin domain; Immunoglobulin V region.  DOMAIN  DOMAIN  98 98 NON_TER 98 98 NON_TER 98 98 AA; 11007 MW; B8644P7F9JFBF95B CRC6	ss-Prot entry is copyright. It is the Swiss Institute of Bioinformation institute. The cong as its content is in no way mo	9654; DOI=10.1084/jem ebecque S.G., Gearhar tion in immunoglobuli 989). g-like (immunoglobuli	1528;  NOV-1990 (Rel. 16, Created)  NOV-1990 (Rel. 16, Last sequence update)  MAY-2005 (Rel. 47, Last annotation update)  heavy chain V region 6.96.  s musculus (Mouse).  karyota; Metazoa; Chordata; Craniata; Verteb  mmalia; Eutheria; Euarchontoglires; Glires;  roidea; Muridae; Murinae; Mus.  1 TaxID-10090;	ALIGNMENTS T 1 MOUSE HV57_MOUSE STANDARD; PRT; 98 AA.	54 58.1 1021 2 Q7SAK8 NEUCR 54 58.1 1037 2 Q6MFQ6_NEUCR 54 58.1 1090 2 Q4NLDB_9MICC 53 57.0 117 1 HV3C_HUMAN 53 57.0 461 2 Q5MFV3_RAT 53 57.0 465 2 Q5MCQ6_HUMAN 53 57.0 465 2 Q6MCQ6_HUMAN 53 57.0 475 2 Q6MCQ6_HUMAN 53 57.0 475 2 Q6MY2_HUMAN 53 57.0 606 2 Q6GMY2_HUMAN 52 56.5 96 2 Q6GMY2_HUMAN 52 56.5 96 2 Q6GMY2_HUMAN 52 55.9 281 2 Q75AG0_ASHGO 52 54.3 116 1 HV05_CARAU 50 53.8 494 2 Q96K68_HUMAN
	Length 98; Indels 0; Gaps 0;	C64;	1 2 2 2	.2007; ; enes du subfami ) domai	rata; Euteleostomi; Rodentia; Sciurognathi;		Q7sak8 neurospora Q6mfq6 neurospora Q4nld8 arthrobacte P01764 homo sapien Q5m7v3 rattus norv Q6p6c4 homo sapien Q6mzq6 homo sapien Q6mzq2 homo sapien Q6my2 homo sapien Q6gmy2 homo sapien Q6zp85 homo sapien Q4x92 tetraodon n Q6zp85 homo sapien Q75ag0 sahbya goss P19181 carassius a P01771 homo sapien

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                                                                                                                                                                               RESULT
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RX MEDLINE=22388257; pubMed=12477932; DOI=10.1073/pnas.242603899;

RX Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

RX Alausner R.D., Collins F.S., Wagner L., Shanmen C.M., Schuler G.D.,

RX Alausner R.D., Collins F.S., Wagner L., Shanefer C.F., Bhat N.K.,

RX Alleschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

RX Alexaner R.D., Collins F.S., Wagner L., Schaefer C.F., Bhat N.K.,

RX Alexaner R.D., Collins F.S., Wagner L., Schaefer C.F., Bhat N.K.,

RX Alexaner R.D., More T., Max S.I., Wang J., Hsieh F.,

RX Alexaner R.D., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

RX Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

RX Alexaner R.D., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

RX Alexaner R.D., Detars G.J., Abramson R.D., Mullahy S.J.,

RX Alexaner R.J., McKernan K.J., Malek J.A., Gunaratne P.H.,

RX Alexaner R.D., Wolffard R.D., More R.D., Mullahy S.J.,

RX Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

RX Villalon D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,

RX Villalon D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,

RX Villalon D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,

RX Alexaner R.D., Dickson M.C.,

RX Alex
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Best Local S
Matches 16
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GO; GO:0003823; F:antigen binding; IEA
InterPro; IPR007110; Ig-like.
InterPro; IPR003597; Ig-cl:
InterPro; IPR003596; Ig-MHC.
InterPro; IPR003996; Ig-W-
Pfam; PP07654; Cl-set; 3.
SMART; SM00406; IGV; 1.
PROSITE; PS50835; IG-LIKE; 4.
PROSITE; PS50835; IG-LIKE; 4.
PROSITE; PS50835; IG-LIKE; 4.
   087 MOUSE
Q32087 MOUSE PRELIMINARY;
Q32087;
Q92087;
01-DEC-2001 (TrembLrel. 19,
01-DEC-2001 (TrembLrel. 19,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Proc. Natl. Acad. Sci [2]
NUCLEOTIDE SEQUENCE.
STRAIN=CZECH II;
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01-JUN-2001 (TrEMBLrel.
01-JUN-2001 (TrEMBLrel.
01-MAR-2004 (TrEMBLrel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TISSUE-Mammary tumor metastatized to Director MGC Project; Submitted (MAR-2001) to the EMBL/GenB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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Name=Igh-VJ558;
Mus musculus (Mouse).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Immunoglobulin domain.
SEQUENCE 487 AA; 57
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EMBL; BC004786; AAH04786.1; -; mRNA.
HSSP; P01810; 2FBJ.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         mouse cDNA sequences.";
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                                                                                                                                                                                                                                                                                                                                                                                                                   l Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; ENSMUSG00000021155; Mus musculus.
                                                                                                                                                                                                                                                                                                                                                 TISDGGSYTYYPDSVKG
          (TrEMBLrel. 19, (TrEMBLrel. 19,
                                                                                                                                                                                                                                                                                                                                                                                                                   96.8%;
llarity 94.1%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Acad. Sci. U.S.A. 99:16899-16903(2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     52555 MW; 7DC8E96DB333077B CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 to the EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             17,
17,
26,
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Last sequence update)
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                                          Created)
          Last sequence update)
                                                                                                                                                                                                                                                                                                                                                                                                                   Score 90; DB 2;
Pred. No. 2.3e-05;
1; Mismatches 0
                                                                                                                PRT;
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RESULT 4

Q652ZL8

AC

Q652ZL8

AC

Q652

D7 25-

D7 25-

D7 25-

D7 25-

D7 Q633

CO Bull

CO Bull

CO Mul

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Best Local S
Matches 16
   PROSITE; PS50835; IGNON TER 196 19 SEQUENCE 196 AA; 2
                                                                                                              InterPro; IPR003599; Ig.
InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig-v.
SMART; SM00409; IG; 1.
SMART; SM00406; IGV; 1.
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Submitted (
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;
                                                                                                                                                                                                                                                                                                                                                                   "An aberrant splicing using a 3' cryptic splice site within exon induces truncated mu-chain production."; Immunology 85:166-170(1995).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NON TER
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InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
SMART; SM00406; IGv; 1.
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PIR; C25913; C25913.
HSSP, P01783; 11GC.
SMR; 092087; 1-119.
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Pterin-mimicking anti-idiotope heavy chain variable region
                                                                                                                                                                                                                                                                                                      EMBL; S79401; AAB35023.2; -; SMR; Q65ZL8; 20-195.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUCLEOTIDE SEQUENCE.
STRAIN=BALB/c;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mus musculus (Mouse)
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Butheria; Euarchontoglires; Glires; Rodentia; Sciurogna
Muridae; Murinae; Mus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQÜENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Komori T., Sugiyama
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDLINE=95362300; PubMed=7635518;
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119 AA;
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                                                 196
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         H.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Euarchontoglires; Glires; Rodentia; Sciurognathi;
               21429 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         93.5%; Score 87; DB 2; 94.1%; Pred. No. 1.5e-0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Jennings I.G., Horaitis O., Cotton R.G.H., to the EMBL/GenBank/DDBJ databases.
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Last sequence update)
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(Fragment).
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               12A381018944B268 CRC64;
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Best Local 9
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InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
SMART; SM00409; IG; 2.
SMART; SM00409; IG; 2.
PROSITE; P850835; IG_LIKE; 2.
NON_TER 1
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05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
                                                                                                                                                                                                                                                                                                    01-NOV-1990 (Rel. 16, Created)
01-NOV-1990 (Rel. 16, Last sequence update)
10-MAY-2005 (Rel. 47, Last annotation update)
1g heavy chain V region 5-76 precursor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Peter J.C., Wallukat G., Tugler J., Maurice D., Roegel J.C., Briand J.P., Hoebeke J.;
"Modulation of the M2 muscarinic acetylcholine receptor activity with monoclonal anti-M2 receptor antibody fragments.";
J. Biol. Chem. 279:55697-55706(2004).
BIOL. GAG34081.1; -; Other_DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eute
Mammalla; Eutheria; Euarchontoglires; Glires; Rodentia;
Muridae; Murinae; Mus.
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Name=scFv B8E5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Q6KB05 MOUSE PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRAIN=Balb/c;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NCBI_TaxID=10090;
MEDLINE-89279149; PubMed=2499654; DOI=10.1084/jem.169.6.2007; Levy N.S., Malipiero U.V., Lebecque S.G., Gearhart P.J.; "Barly onset of somatic mutation in immunoglobulin VH genes d primary immune response."; J. Exp. Med. 169:2007-2019(1989).
                                                                                                                                                                       NCBI_TaxID=10090;
                                                                                                                                                                                                              Mus musculus (Mouse)
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;
Muroidea; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                         HV58 MOUSE
P18529;
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                                                                                                                      NUCLEOTIDE SEQUENCE.
STRAIN=BALB/cJ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            255 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            27445 MW;
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Pred. No.
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Pred. No. (
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tia; Sciurognathi;
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PDB; 1181; X-ray; B=26-115.

PDB; 1188; X-ray; B=26-115.

PDB; 1188; X-ray; B=26-115.

InterPro; IPR007710; Ig-like.

InterPro; IPR007516; Ig_v:

SMART; SM00406; IGv; II_v:

SMART; SM00406; IGv; I.

PROSITIE; PS50835; IG_LIKE; 1.

3D-structure; Immunoglobulin do SIGNAL 1 19

CHAIN 20 197

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P18530;
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                                                                                                                                                                   PIR; JT0507; HVMS39.
HSSP; P18529; 118K.
SMR; P18530; 20-117.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 primary immune response.";
J. Exp. Med. 169:2007-2019(1989).
-!- MISCELLANEOUS: This sequence belongs to the VH7183 subfamily
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=89279149; PubMed=2499654; DOI=10.1084/jem.169.6.2007;
Levy N.S., Malipiero U.V., Lebecque S.G., Gearhart P.J.;
"Early onset of somatic mutation in immunoglobulin VH genes of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUCLEOTIDE SEQUENCE.
STRAIN=BALB/cJ;
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Mammalia, Eutheria, Euarchontoglires, Glires, Rodentia, Sciurognathi,
Muroidea, Muridae, Murinae, Mus.
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                                                                                                                                                                                                                                                                                                                                                          use
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88.2%;
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Pred. No. 0.00013;
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SMART; SM00406; IGV; 1. PROSITE; PS50835; IG\_LIKE; 1.

Immunoglobulin V region;

InterPro; IPR007110; Ig-like. InterPro; IPR003596; Ig\_v.

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01-NOV-1990 (Rel. 16, Last sequence update)
10-MAY-2090 (Rel. 16, Last sequence update)
10-MAY-2090 (Rel. 47, Last annotation update)
Ig heavy chain V region 5-84 precursor.
Mus musculus (Mouse).
Mus musculus (Mouse).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Euarchontoglires, Glires, Rodentia, Sciurognathi,
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01-NOV-1990
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InterPro; IPR003596; Ig_v.
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J. Exp. Med. 169:2007-2019(1989).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Levy N.S., Malipiero U.V., Lebecque S.G., Gearhart P.J.; Barly onset of somatic mutation in immunoglobulin VH genes during
                                                                                                                                                                                                                                                                                                                   3D-structure; Immunoglobulin
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1AR1; X-ray; D=-
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ISDGGSYTYYPDSVKG 17
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82.4%;
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75.0%;
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Framework-1.
Complementarity-determining-1.
Framework-2.
Complementarity-determining-2.
Framework-3.
By similarity.
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Pred. No.
                                                         Score 67; DB 1;
Pred. No. 0.02;
                                                                                                                                                      Framework-3.
By similarit
                                                                                                                                                                                     Framework-2.
Complementarity-determining-2.
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Complementarity-determining-1.
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RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

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RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

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RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

RA Raha S.S., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,

RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

RA Whiting M., Madan A., Young A.C., Shevchbako Y., Bouffard G.G.,

RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,

RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,

RA Schnerch A., Schein J.E., Jones S.J.M., Marra M.A.,

"Generation and initial analysis of more than 15,000 full-length human

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InterPro; IPR007110; Ig-like.
InterPro; IPR003597; Ig_cl.
InterPro; IPR003006; Ig_MHC.
InterPro; IPR003596; Ig_v.
Pfam; PP07654; Cl-set; 3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Klein S., Gerhard D.S.;
Submitted (OCT-2004) to the EMBL/GenBank/DDBJ databases.
EMBL; BC084123; AAH84123.1; -; mRNA.
GO; GO:0030106; F:MHC class I receptor activity; IEA.
GO; GO:0019883; P:antigen presentation, endogenous antigen; I
GO; GO:0019885; P:antigen processing, endogenous antigen via
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Mesobatrachia; Pipoidea; Pipidae;
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Proc. Natl. Acad. Sci. U.S.A.
                                                                                            PROSITE; PS50835; IG_LIKE; 5.
PROSITE; PS00290; IG_MHC; UNKNOWN_4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
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RESULT 10

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RESULT 11
HV56 MOUSE STANDARD;
ID HV56 MOUSE STANDARD;
AC P18527;
DT 01-NOV-1990 (Rel. 16, Created)
DT 01-NOV-1990 (Rel. 16, Last seq
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Matches 12
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Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurogna
Muroidea; Muridae; Murinae; Mus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ig heavy chain V region Mus musculus (Mouse).
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10-MAY-2005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This Swiss-Prot entry is copyright. It is produced through a collabout between the Swiss Institute of Bioinformatics and the EMBL outstathe European Bioinformatics Institute. There are no restrictions cure as long as its content is in no way modified and this statement
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          primary immune response.";
J. Exp. Med. 169:2007-2019(1989).
-!- MISCELLANEOUS: This sequence belongs to the VH7183 subfamily.
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InterPro; IPR0077110; Ig-11ke.
InterPro; IPR003596; Ig-v.
SMART; SM00406; IGv; 1.
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Pred. No.
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Pred. No.
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By similarity.
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Complementarity-determining-2.
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Mammalia; Eutheria; Euarchontoglires; Glires;
Muroldea; Muridae; Murinae; Mus.
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Levy N.S., Malipiero U.V., Lebecque S.G., Gearhart P.J.;
"Early onset of somatic mutation in immunoglobulin VH genes during
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NON TER
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NUCLEOTIDE SEQUENCE.

NUCLEOTIDE SEQUENCE.

Jailon O., Aury J.M., Brunet F., Petit J.L., Stange-Thomann N.,

Jailon O., Aury J.M., Brunet F., Cozouf-Costaz C., Bernot A.,

Maucell E., Bouneau L., Pischer C., Ozouf-Costaz C., Bernot A.,

Nicaud S., Jaffe D., Fisher S., Lutfalla G., Dossat C., Segurens B.

Dasilva C., Salanoubat M., Levy M., Boudet N., Castellano S.,

Anthouard V., Jubin C., Castelli V., Katinka M., Vacherie B.,

Biemont C., Skalli Z., Cattolico L., Poulain J., De Berardinis V.,

Cruaud C., Duprat S., Brottier P., Coutanceau J.P., Gouzy J.,

Parra G., Lardier G., Chapple C., McKernan K.J., McEwan P., Bosak &

Kellis M., Volff JN., Guigo R., Zody M.C., Mesirov J.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORFNames=GSTENG00004372001;
Tetraodon nigroviridis (Green puffer).
Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Eukaryota; Metazoa; Chordata; Teleostei; Euteleostei; Neoteleostei;
Actinopterygii; Teleostei; Euteleostei; Neoteleostei;
Acanthomorpha; Acanthopterygii; Percomorpha; Tetraodontiformes;
Tetradontoidea; Tetraodontidae; Tetraodon.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            13-SEP-2005 (TrEMBLrel. 31, Created)
13-SEP-2005 (TrEMBLrel. 31, Last sequer
13-SEP-2005 (TrEMBLrel. 31, Last annote
Chromosome undetermined SCAF649, whole
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Q4TJ56;
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SMR; P18527; 1-97.
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82.4%;
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whole genome shotgun sequence.
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A Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schular G.D.,
A Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
A Altschul S.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
A Hopkins R.P., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
A Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
A Diatchenko L., Warusina K., Farmer A.A., Rubin G.M., Hong L.,
A Diatchenko L., Warusina K., Farmer A.A., Rubin G.M., Hong L.,
A Diatchenko L., Warusina K., Farmer A.A., Rubin G.M., Hong L.,
A Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
A Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
A Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
A Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
Yillalon D.K., Mozny D.M., Sodergren B.J., Lu X., Gibbs R.A.,
Yallalon D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,
Yallalon D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,
Yallalon D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,
Yallalon D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,
Yallalon D.K., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
A Whiting M., Wadan A., Young A.C., Schwutz J., Myers R.M.,
A Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
A Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
Yallalon J.E., Jones S.J.M., Marra M.A.;
"Generation and initial analysis of more than 15,000 full-length human
"Tend mouse Chub."
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                                  NIH MGC Project;
Submitted (MAR-2005) to the
EMBL; BC091247; AAH91247.1;
SMR; Q5EK12; 20-241.
GO; GO:0003823; F:antigen b:
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;
Muroidea; Muridae; Murinae; Rattus.
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-i- CAUTION: The sequence shown here is derived from an EMBL/GenBank/DDBJ whole genome shotgun (WGS) entry whi
                                                                                                                                                                                         NUCLEOTIDE SEQ
TISSUE=Spleen;
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j; CAAE01000649; CAF87076.1; -; Genomic_DNA.
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A Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
A Strausberg R.L., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
A Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Heich F.,
A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Heich F.,
A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Heich F.,
A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Heich F.,
A Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
A Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
A Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
A Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
A Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
A Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
A Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
A Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
A Willalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
A Willalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
A William M., Madan A., Young A.C., Shevbneko Y., Bouffard G.G.,
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InterPro; IPR003597; Ig c1.
InterPro; IPR003597; Ig MHC.
InterPro; IPR003596; Ig_v.
Pfam; PP07554; C1-set; 2.
SMART; SM00409; IG; 3.
SMART; SM00409; IGc1; 3.
SMART; SM00406; IGv; 1.
PROSITE; P850835; IG LIKE; 3.
PROSITE; P850835; IG LIKE; 3.
PROSITE; P850290; IG MHC; UNKNOWN 2
SEQUENCE 479 AA; 52329 MW; 8BI44
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13-SEP-2005 (TrEMBLrel. 3
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Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C., Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.N., Krzywinski M.I., Skalska U., Smailus D.B. Schmerch A., Schein J.E., Jones S.J.M., Marra M.A.; Scherch A., Schein J.E., Jones S.J.M., Marra M.A.; "Generation and initial analysis of more than 15,000 full-len and mouse cDNA sequences."; proc. Natl. Acad. Sci. U.S.A. 99:16899-16903 (2002).
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Mammalia, Eutheria, Euarchontoglires, Glires, Rodentia, Sciurogna
Muroidea, Muridae, Murinae, Rattus.
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DT 05-JUL-2004 (TREMBLIE1. 27, 1)
DE HYPOTHBEIGE (Human).

CC HOMO SAPIERS (Human).
CC EUKARYOTA; MCTEACA; CHOOTOCATA
CC EUKARYOTA; SEQUENCE.
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Search completed: November 21, 2005, 12:04:11 Job time: 26.1575 secs
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R HSSP; P01861; 1ADQ.

R InterPro; IPR003599; Ig.

R InterPro; IPR007110; Ig-like.

InterPro; IPR003197; Ig-like.

InterPro; IPR003006; Ig-MHC.

R InterPro; IPR003596; Ig-WHC.

R InterPro; IPR003596; Ig-WHC.

R InterPro; IPR003596; Ig-WHC.

R InterPro; IPR003596; Ig-WHC.

R Pfam; PF07654; C1-8et; 3.

R SMART; SM00409; IG; 2.

R SMART; SM00409; IG; 1.

R SMART; SM00406; IGv; 1.

R PROSITE; PS0033; IG LIKE; 4.

PROSITE; PS00339; IG LIKE; 4.

Hypothetical protein.

O SEQUENCE 464 AA; 51099 MW; 2FCA72C66EBA0ABC CRC64;
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SMART; SM00407; IGcl; 2.
SMART; SM00406; IGcl; 2.
PROSITE; PS50835; IG_LIKE; 4.
Immunoglobulin domain; Repeat.
SEQUENCE 467 AA; 51651 MW; 1FF0328F50160ED3 CRC64;
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05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
Hypothetical protein DKPZp686C15213.
Name=DKFZp686C15213;
Name=DKFZp686C15213;
Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Bukaryota; Metazoa; Chordata; Craniata; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The German cDNA Consortium;
Bloecker H., Beccher M., Brandt P., Mewes H.W., Weil B.,
Osanger A., Pobo G., Han M., Wiemann S.;
Submitted (JAN-2005) to the EMBL/GenBank/DDBJ databases.
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score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.
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Copyright (c) 1993 - 2005 Compugen Ltd.
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| hes 17           | AT TE TE TE O TE TE TE TE TE TE TE TE TE TE TE TE TE | TITLE OF INVENTION: HUMA TITLE OF INVENTION: IMMU NUMBER OF SEQUENCES: 28 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy di COMPUTER: IBM PC COMPO OPERATING SYSTEM: PC-D SOFTWARE: ASCII text CURRENT APPLICATION UMBER: US FILING DATE: CHASSIFICATION: 424 PRIOR APPLICATION UMBER: US APPLICATION UMBER: US FILING DATE: CHASSIFICATION UMBER: US APPLICATION UMBER: US APPLICATION UMBER: US FILING DATE: 04-NOV-19 APPLICATION UMBER: US                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    | IT 1<br>1-435-516-<br>puence 1,<br>puence 1,<br>NERAL ING. 6 | 28 87 30 87 31 87 32 87 34 87 35 87 36 87 36 87 36 87 36 87 36 87 36 87 87 87 87 87 87 87 87 87 87 87 87 87                                                                                                                                                                                                                                                                                                                                                                                                                      |
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| Conse            | IN IN IN IN IN IN IN IN IN IN IN IN IN I             | TIE OF INVENTION: HUTLE OF INVENTION: HUTLE OF INVENTION: IMPORTER READABLE FOOM: MEDIUM TYPE: Floppy MEDIUM TYPE: Floppy MEDIUM TYPE: Floppy MEDIUM TYPE: POOM: MEDIUM TYPE: POOM: MEDIUM TYPE: POOM: MEDIUM TYPE: POOM: MEDIUM TYPE: POOM: MEDIUM TYPE: POOM: MEDIUM TYPE: POOM: MEDIUM TYPE: POOM: MEDIUM TYPE: POOM: MEDIUM TYPE: MEDIUM TYPE: MEDIUM TOWN MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM MEDIUM M | ion                                                          | 93.5<br>93.5<br>11<br>93.5<br>13<br>93.5<br>13<br>93.5<br>14<br>93.5<br>14<br>93.5<br>14<br>93.5<br>14<br>93.5<br>14<br>93.5<br>14<br>93.5<br>14<br>93.5<br>14<br>93.5<br>14<br>93.5<br>14<br>93.5<br>14<br>93.5<br>14<br>93.5<br>14<br>93.5<br>14<br>93.5<br>14<br>93.5<br>14<br>93.5<br>14<br>93.5<br>14<br>93.5<br>14<br>93.5<br>14<br>93.5<br>14<br>93.5<br>14<br>93.5<br>14<br>93.5<br>14<br>93.5<br>14<br>93.5<br>14<br>93.5<br>14<br>93.5<br>14<br>93.5<br>15<br>16<br>16<br>16<br>16<br>16<br>16<br>16<br>16<br>16<br>16 |
|                  | N: 5,4                                               | ON: HUMANIZED A ON: IMMUNOBLOBI CES: 28 E FORM: E FORM: PC Compatible PC Compatible II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II text II | A<br>US/08435516                                             | 8887763556563419999<br>666666666666666666666666666666666                                                                                                                                                                                                                                                                                                                                                                                                                                                                         |
| 0; Mismatches 0  | N:<br>N: 013<br>N: DB 2;<br>Pred. No. 7.5e-07;       | ED ANTIBODIÉS TO POLOBULIN G ON HUMAN  Le MS-DOS  /435,516  23377.4; PCT/US93,-02-NOV-1993                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     | ALIGNMENTS                                                   | US-09-269-332-86 US-09-269-332-90 US-09-269-332-77 US-09-269-332-77 US-09-269-628-23 US-09-069-628-23 US-09-069-628-23 US-09-069-628-25 US-09-840-459-45 US-09-840-459-45 US-09-495-880A-11 US-08-356-272-3 US-09-495-880A-11 US-08-356-272-3 US-09-318-737-2 US-09-69-628-25 US-09-69-628-25 US-09-69-628-25 US-09-69-628-25 US-09-69-674A-1 US-08-875-674A-3 US-08-875-674A-3 US-08-875-674A-3 US-08-875-674A-3                                                                                                                |
| Indels 0; Gaps 0 | Length 120;                                          | RECEPTORS FOR PHAGOCYTES                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       |                                                              | Sequence 56, Appl<br>Sequence 88, Appl<br>Sequence 90, Appl<br>Sequence 76, Appl<br>Sequence 77, Appl<br>Sequence 23, Appl<br>Sequence 26, Appl<br>Sequence 45, Appl<br>Sequence 45, Appl<br>Sequence 42, Appl<br>Sequence 11, Appl<br>Sequence 27, Appli<br>Sequence 27, Appli                      |

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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: ASCII text
CURRENT APPLICATION DATA:
   atent No. 6500931
   equence 3, Application US/08435516 atent No. 6500931 GENERAL INFORMATION:
APPLICANT:
   REGISTRATION NUMBER: 35,430
REFERENCE/DOCKET NUMBER: MX:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 227-7400
TELEFAX: (617) 227-5941
INFORMATION FOR SEQ ID NO: 2:
   BPPLICANT:
REGISTRATION NUMBER: 35,430
REFERENCE/DOCKET NUMBER: MXI-013
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 227-7400
  ATTORNEY/AGENT INFORMATION:
NAME: Arnold, Beth E.
   PRIOR APPLICATION DATA:
APPLICATION NUMBER: UK 92 23377.4; PCT/US93/10384
FILING DATE: 04-NOV-1992; -02-NOV-1993
CLASSIFICATION: 424
  NUMBER OF SEQUENCES: 2
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MEDIUM TYPE: Floppy
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   MOLECULE TYPE: peptide FRAGMENT TYPE: internal
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LENGTH: 120 amino acid
  ATTORNEY/AGENT INFORMATION:
NAME: Arnold, Beth E.
   PRIOR APPLICATION DATA:
APPLICATION NUMBER: UK 92 23377.4; PCT/US93/10384
FILING DATE: 04-NOV-1992; -02-NOV-1993
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  FILING DATE:
CLASSIFICATION:
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  1 TISDGGSYTYYPDSVKG 17
   TISDGGSYTYYPDSVKG 66
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  Conservative
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   HUMANIZED ANTIBODIES TO FC RECEPTORS FOR IMMUNOBLOBULIN G ON HUMAN MONONUCLEAR PHAGOCYTES
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Pred. No. 7.5e-07;
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  片
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US-08-661-052-4
  US-09-344-587-13
  US-09-344-587-13
   US-08-435-516-3
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CURRENT FILING DATE: 1999-06-25
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EARLIER FILING DATE: 1998-06-26
NUMBER OF SEQ ID NOS: 14
SOFTWARE: Patentin Ver. 2.1
  INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS
   GENERAL INFORMATION:
APPLICANT: Yashwant M. Deo
APPLICANT: Joel Goldstein
APPLICANT: Bobert Graziano
APPLICANT: Chezian Somasundaram
APPLICANT: Chezian Somasundaram
TITLE OF INVENTION: THERAPBUTIC COMPOUNDS COMPRISED
TITLE OF INVENTION: OF ANTI-FC RECEPTOR ANTIBODIES
  TITLE OF INVENTION: REDUCTION OF INTERFERENCE OF IMMUNOASSAYS BY SUBSTANCES TITLE OF INVENTION: DERIVED FROM THE FRAMEWORK REGIONS OF ANTIBODIES FILE REFERENCE: BMID 9928 4897/00/US-Im
   APPLICANT: Praast, Gerald APPLICANT: Nussbaum, Sabine APPLICANT: Moessner, Ellen APPLICANT: Lenz, Helmut
  ORGANISM: Mus musculus
  TYPE: PRT
  LENGTH: 123
ZIP: 02109-1875
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  TOPOLOGY:
  ADDRESSEE:
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  1 TISDGGSYTYYPDSVKG 17
   1 TISDGGSYTYYPDSVKG 17
  Boston
   Similarity
  amino acid
  Massachusetts
  Application US/09344587
  60 State Street, Suite 510
  120 amino acids
   USA
   Conservative
  LAHIVE & COCKFIELD
   100.0%; Score 93; DB 2; 100.0%; Pred. No. 7.7e-07;
  ω
••
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  Matches
   Query Match
  PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/
FILING DATE: 07-JUNE-1995
ATTORNEY/AGENT INFORMATION:
NAME: ATTOId, Beth E.
   GENERAL INFORMATION:
  TELEFAX: (617)227-5941 (NFORMATION FOR SEQ ID NO: 4:
   APPLICATION NUMBER: U:
FILING DATE:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08
FILING DATE:
             TELEPAX: (617)227-5941
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
  SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
  APPLICANT:
APPLICANT:
APPLICANT:
  MOLECULE TYPE: peptide FRAGMENT TYPE: internal
   REFERENCE/DOCKET NUMBER: MX
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)227-7400
  SEQUENCE CHARACTERISTICS:
  ZIP: 02109-1875
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
   COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
   APPLICANT: Chezian Somasundaram
TITLE OF INVENTION: THERAPEUTIC COMPOUNDS COMPRISED
TITLE OF INVENTION: OF ANTI-FC RECEPTOR ANTIBODIES
   CORRESPONDENCE ADDRESS:
   NUMBER OF SEQUENCES:
  NAME: Arnold, Beth E.
REGISTRATION NUMBER: 35,430
REFERENCE/DOCKET NUMBER: WXI-043CP
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)227-7400
   ATTORNEY/AGENT INFORMATION:
  y Match 100.0%;
Local Similarity 100.0%;
hes 17; Conservative (
   LENGTH: 300 amino acids
TYPE: amino acid
TOPOLOGY: linear
   REGISTRATION NUMBER: 35,430
   COUNTRY:
  CITY: Boston
  STREET:
   FILING DATE:
  ADDRESSEE: LAHIVE & COCKFIELD
   50
  1 TISDGGSYTYYPDSVKG 17
   TISDGGSYTYYPDSVKG 66
   Massachusetts
  60 State Street, Suite 510
  Joel Goldstein
Robert Graziano
   USA
300 amino acids
   Yashwant M. Deo
  08/661,052
   US/09/188,082
   08/484,172
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   MXI-043CP
   Score 93; DB 1;
Pred. No. 2e-06;
); Mismatches
   DB 1; Length 300;
   Indels
   0
   Gaps
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   S
  US-09-188-082-4
  RESULT 7
US-09-364-088-4
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밁
  US-09-364-088-4
   REFERENCE/DOCKET NUMBER: MXI-
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)227-7400
TELEPHAX: (617)742-77414
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 300 amino acids
  Patent No.
   Sequence 4, Application US/09364088
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   Query Match
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Best Local Similarity
Matches 17; Conserv
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APPLICANT: Yashwa:
   OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIA Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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  TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
   APPLICATION NUMBER: US 08/484,172
FILING DATE: 07-JUNE-1995
ATTORNEY/AGENT INFORMATION:
NAME: 07-111
   COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
  NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
   TITLE OF INVENTION: THERAPEUTIC COMPOUNDS COMPRISE TITLE OF INVENTION: OF ANTI-FC RECEPTOR ANTIBODIES
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FILING DATE: 07-JUNE-1996
PRIOR APPLICATION DATA:
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TOPOLOGY: linear
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   STREET:
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   1 TISDGGSYTYYPDSVKG 17
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TISDGGSYTYYPDSVKG 66
                   TISDGGSYTYYPDSVKG 17
  Remillard, Jane
   Massachusetts
   E: LAHIVE & COCKFIELD, LLP
28 State Street, 24th Floor
  100.0%; ilarity 100.0%; Conservative 0;
  Yashwant M. Deo, et al. // THERAPEUTIC COMPOUNDS COMPRISED
   Floppy disk
   09/188,082
   66
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  MXI-043CP2
   Score 93; DB 2;
Pred. No. 2e-06;
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RESULT 8 US-09-102-716-4

Sequence 4, Application US/09102716

Patent No. 6395272
GENERAL INFORMATION:

APPLICANT: Yashwant M. Deo

Joel Goldstein Robert Graziano Chezian Somasundaram

TITLE OF INVENTION: THERAPEUTIC COMPOUNDS COMPRISED

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US-08-661-052-14
  Matches
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  GENERAL INFORMATION:
APPLICANT: Yashwant M. Deo
APPLICANT: Joel Goldstein
APPLICANT: Robert Graziano
APPLICANT: Chezian Somasund
  TELEFAX: (617)227-5941
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
   tent No.
   APPLICANT: Chezian Somasundaram
TITLE OF INVENTION: THERAPEUTIC COMPOUNDS COMPRISED
TITLE OF INVENTION: OF ANTI-FC RECEPTOR ANTIBODIES
NUMBER OF SEQUENCES: 16
CORRESPONDENCE ADDRESS:
ADDRESSES: LAHIVE & COCKFIELD
STREET: 60 State Street, Suite 510
   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION NUMBER: US/09/102,716
APPLICATION NUMBER: US/09/102,716
FILING DATE: 22-un-1998
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/661,052
PILING DATE: <Unknown>
COMPUTER READABLE FORM:
   STREET: 60 State Stre
CITY: Boston
STATE: Massachusetts
  Local
                COUNTRY: USA
ZIP: 02109-1875
   MOLECULE TYPE: peptide FRAGMENT TYPE: internal SEQUENCE DESCRIPTION: SEQ ID NO: 4:
  NAME: Arnold, Beth E.
REGISTRATION NUMBER: 35,430
REFERENCE/DOCKET NUMBER: MXI-043CP
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)227-7400
  NUMBER OF SEQUENCES: 16
CORRESPONDENCE ADDRESS:
ADDRESSEE: LAHIVE & COCKFIELD
  ATTORNEY/AGENT INFORMATION:
   50
  1 TISDGGSYTYYPDSVKG 17
  4, Application US/08661052
5837243
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   TISDGGSYTYYPDSVKG 66
   TYPE: amino acid
  COUNTRY: USA
ZIP: 02109-1875
  STATE: Massachusetts
  CITY: Boston
   STREET: 60 State Street, Suite 510
   TOPOLOGY: linear
   LENGTH: 300 amino acids
   100.0%; Score 93; DB 2; Length 300; 100.0%; Pred. No. 2e-06; tive 0; Mismatches 0; Indels
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RESULT 10
US-09-188-082-14
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  Sequence 14, Application US/09188082 Patent No. 6270765
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               TELEPAX: (617)227-594
INFORMATION FOR SEQ ID NO:
  TELEFAX: (617)227-5941
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
  GENERAL INFORMATION:
   MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/661,052
   NAME: Arnold, Beth E.
REGISTRATION NUMBER: 35,430
REFERENCE/DOCKET NUMBER: MX.
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)227-7400
  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
  FILING DATE: 07-JUNE-1995
ATTORNEY/AGENT INFORMATION:
NAME: Arnold, Beth E.
   PILING DATE:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
   APPLICANT: Yashwant M. Deo
APPLICANT: Joel Goldstein
APPLICANT: Robert Graziano
APPLICANT: Chezian Somasundaram
   REGISTRATION NUMBER: 35,430
REFERENCE/DOCKET NUMBER: MX
TELECOMMUNICATION INFORMATION:
SEQUENCE CHARACTERISTICS:
  FILING DATE:
ATTORNEY/AGENT INFORMATION:
   PRIOR APPLICATION DATA:
APPLICATION NUMBER:
  NUMBER OF SEQUENCES: 16
CORRESPONDENCE ADDRESS:
ADDRESSEE: LAHIVE & COCKFIELD
   TITLE OF INVENTION: THERAPEUTIC COMPOUNDS COMPRISED TITLE OF INVENTION: OF ANTI-FC RECEPTOR ANTIBODIES
   APPLICATION NUMBER: FILING DATE:
   Natch 100.0%; Score 93; Local Similarity 100.0%; Pred. No.
  TYPE: amino acid
TOPOLOGY: linear
  STREET:
   COUNTRY:
   TELEPHONE:
  REGISTRATION NUMBER:
  LENGTH: 301 amino acids
  69 TISDGGSYTYYPDSVKG
  1 TISDGGSYTYYPDSVKG 17
   02109-1875
  Boston
   Massachusetts
  60 State Street, Suite 510
   (617)227-5941
TO NO: 14:
   USA
  Conservative
                                 (617) 227-5941
   S
   08/661,052
   US/09/188,082
   08/484,172
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  MXI-043CP
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  Gaps
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The second state

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WS-09-102-716-14
US-09-102-716-14
; Sequence 14, Application US/09102716
; Patent No. 6395272
   ; TOPOLOGY: linear; MOLECULE TYPE: protein US-09-364-088-14
   ATTORNEY/AGENT INFORMATION:
NAME: Remillard, Jane E.
REGISTRATION NUMBER: 38,872
REFERENCE/DOCKET NUMBER: MXI-043CP2
REFERENCE/INFORMATION:
TELEPHONE: (617)227-7400
TELEPAX: (617)742-7414
INFORMATION FOR SEQ ID NO: 14:
   Query Match 100.0%;
Best Local Similarity 100.0%;
Matches 17; Conservative 0
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Best Local Similarity
Matches 17; Conserv
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  APPLICATION NUMBER: US/09/364,088
FILLING DATE:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 09/188,082
FILLING DATE: 07-JUNE-1996
PRIOR APPLICATION NUMBER: US 08/484,172
APPLICATION NUMBER: US 08/484,172
FILLING DATE: 07-JUNE-1995
  GENERAL INFORMATION:

APPLICANT: Yashwant M. Deo, et al.

TITLE OF INVENTION: THERAPEUTIC COMPOUNDS COMPRISED

TITLE OF INVENTION: OF ANTI-FC RECEPTOR ANTIBODIES
  tent No.
  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: DM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
   NUMBER OF SEQUENCES: 16

CORRESPONDENCE ADDRESS:

ADDRESSEE: LAHIVE & COCKFIELD, LLP
STREET: 28 State Street, 24th Floor
  SEQUENCE CHARACTERISTICS:
LENGTH: 301 amino acids
  TYPE: amino acid
TOPOLOGY: linear
   LENGTH:
   TYPE: amino acid
   COUNTRY:
  CITY: Boston
   69
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   09/188,082
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Pred. No. 2.1e-06;
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APPLICANT: GENER, A.O.

APPLICANT: GRAZIANO, Robert

APPLICANT: DEO, YASHWANT M.

TITLE OF INVENTION: CELLS EXPRESSING ANTI-FC RECEPTOR

TITLE OF INVENTION: BINDING COMPONENTS

FILE REFERENCE: MXI-099CPA

CURRENT APPLICATION NUMBER: US/09/203,958A

CURRENT TILING DATE: 1998-12-02

PRIOR APPLICATION NUMBER: 60/067232

PRIOR FILING DATE: 1997-12-02

INVMER OF SEQ ID NOS: 4

SOPTWARE: FASTSEQ for Windows Version 4.0

SEQ ID NO 2

SEQ ID NO 2
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US-09-203-958A-2
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   GENERAL INFORMATION:
APPLICANT: KELER, T
APPLICANT: GOLDSTE
   Sequence 2, Application US/09203958A Patent No. 6682928
   Matches
  Query Match 100.0%;
Best Local Similarity 100.0%;
  TELEPHONE: (617)227-74
TELEPAX: (617)227-5941
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MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/102,716

PILING DATE: 22-Jun-1998

PRIOR APPLICATION NUMBER: 08/661,052

PILING DATE: <Unknown>

APPLICATION NUMBER: 08/661,052

PILING DATE: <Unknown>

APPLICATION NUMBER: 35,430

REFERENCE/DOCKET NUMBER: MXI-043CP

TELEGOMMUNICATION INFORMATION:

TELEGOMMUNICATION INFORMATION:
  LENGTH: 301 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
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  CORRESPONDENCE ADDRESS:
  NUMBER OF SEQUENCES: 16
  Chezian Somasundaram
TITLE OF INVENTION: THERAPEUTIC COMPOUNDS COMPRISED
OF ANTI-FC RECEPTOR ANTIBODIES
   APPLICANT: Yashwant M. Deo
  69
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  COUNTRY: USA
   STATE: Massachusetts
  CITY: Boston
   STREET: 60 State Street, Suite 510
  DDRESSEE: LAHIVE & COCKFIELD
  Robert Graziano
   Joel Goldstein
  Tibor
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Pred. No. 2.1e-06;
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  Gaps
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Sequence 16, Application US/09188082
Patent No. 6270765
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APPLICANT: Yashwant M. Deo
  Matches
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Best Local (
   GENERAL INFORMATION:

APPLICANT: Yashwant M. Deo
APPLICANT: Joel Goldstein
APPLICANT: Robert Graziano
APPLICANT: Chezian Somasundaram
TITLE OF INVENTION: THERAPEUTIC COMPOUNDS COMPRISED
TITLE OF INVENTION: OF ANTI-FC RECEPTOR ANTIBODIES
NUMBER OF SEQUENCES: 16
  APPLICATION NUMBER: US/08/661
FILING DATE:
PRIOR APPLICATION DATA:
APPLICATION UMBER: US 08/48
APPLICATION NUMBER: US 08/48
FILING DATE: 07-JUNE-1995
ATTORNEY/AGENT INFORMATION:
NAME: ARROLD, Beth E.
REGISTRATION NUMBER: 35,430
REFERENCE/DOCKET NUMBER: MXI-
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
  TELEFAX: (617)227-5941
INFORMATION FOR SEQ ID NO: 16:
SEQUENCE CHARACTERISTICS:
LENGTH: 553 amino acids
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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ADDRESSEE: LAHIVE & COCKFIELD
STREET: 60 State Street, Suite 510
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밁 Ś

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  TELEPHONE: (617)227-7400
TELEFAX: (617)227-5941
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TYPE: amino acid
TOPOLOGY: linear
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  Query Match
  FILING DATE:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
  ATTORNEY/AGENT INFORMATION:
NAME: Arnold, Beth E.
REGISTRATION NUMBER: 35,430
REFERENCE/DOCKET NUMBER: MX:
TELECOMMUNICATION INFORMATION:
  COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
  NUMBER OF SEQUENCES: 16
CORRESPONDENCE ADDRESS:
ADDRESSEE: LAHIVE & COCKFIELD
STREET: 60 State Street, Suite 510
  APPLICANT: Joel Goldstein
APPLICANT: Robert Graziano
APPLICANT: Chezian Somasundaram
TITLE OF INVENTION: THERAPEUTIC COMPOUNDS COMPRISED
TITLE OF INVENTION: OF ANTI-FC RECEPTOR ANTIBODIES
   / Match 100.0%; Score 93; DB 2; Length 553; Local Similarity 100.0%; Pred. No. 4e-06;
   CITY: Boston
  APPLICATION NUMBER: US/09/188,082
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  Mismatches
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  1 TISDGGSYTYYPDSVKG 17
Published Applications AA Main:*

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2: /cgn2 6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

3: /cgn2 6/ptodata/1/pubpaa/US09_PUBCOMB.pep:*

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  GenCore version 5.1.6 Copyright (c) 1993 - 2005 Compugen Ltd.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

| 27                | 26             | 25                | 24           | 23             | 22            | 21               | 20    | 19            | 18                | 17            | 16           | 15           | 14            | 13           | 12              | 11              | 10              | 9                | 80    | 7                | 6                | <sub>5</sub>      | 4                 | w                 | ຎ      | ь                 | Result           |
|-------------------|----------------|-------------------|--------------|----------------|---------------|------------------|-------|---------------|-------------------|---------------|--------------|--------------|---------------|--------------|-----------------|-----------------|-----------------|------------------|-------|------------------|------------------|-------------------|-------------------|-------------------|--------|-------------------|------------------|
| 87                | 87             | 87                | 87           | 90             | 90            | 93               | 93    | 93            | 93                | 93            | 93           | 93           | 93            | 93           | 93              | 93              | 93              | 93               | 93    | 93               | 93               | 93                | 93                | 93                | 93     | 93                | Score            |
| 93.5              | ٠              | 93.5              | •            | ٠              | •             | •                | •     | •             | •                 | 100.0         | •            | 100.0        | 100.0         | •            | 100.0           | •               | •               | •                | 100.0 | ٥.               | •                | 100.0             |                   | •                 | 100.0  | 100.0             | Query<br>Match 1 |
| 17                | 17             | 17                | 17           | 123            | 123           | 4852             | 663   | 352           | 352               | 120           | 120          | 120          | 120           | 120          | 120             | 120             | 120             | 118              | 118   | 118              | 118              | 17                | 17                | 17                | 17     | 17                | Length [         |
| 4.                | 4              | Δ.                | 4.           | 4              | w             | 4                | 4     | v             | w                 | 4.            | 4            | 4            | 4             | 4            | 4               | 4               | 4               | 4                | 4.    | w                | w                | 4                 | 4                 | 4                 | 4      | 4.                | . B              |
| US-10-477-377-9   | -10-286-132A-2 | -10-275-180A-     | 0-281-479A-2 | -10-632-706-57 | -09-144-886-6 | US-10-412-406-33 | ដំ    | -10-764-131-2 | -09-203-958A      | -10-412-406-1 | -10-412-406- | -10-412-406- | -10-412-406-1 | -10-412-406- | US-10-229-335-3 | US-10-229-335-2 | US-10-229-335-1 | US-10-632-706-60 | -706- | US-09-144-886-63 | US-09-144-886-62 | US-10-632-706-157 | US-10-632-706-153 | US-10-632-706-149 |        | US-10-632-706-141 | Ĭ                |
| sequence 9, Appli | 6              | Sequence 26, Appl | 26,          | e 57,          | 60            | e 33,            | e 32, | e 2,          | Sequence 2, Appli | e 16          | e 14         | e 12         | e 10,         | e 2, 1       | e<br>3,         | е 2,            | e 1, 1          |                  | e 59, | e 63,            | •                | e 157,            | e 153,            | e 149,            | e 145, | e 141,            | Description      |

| 45               | 44               | 43               | 42               | 41               | 40               | 39               | 38               | 37               | 36               | 35               | 34               | 33               | 32               | 31               | 30               | 29               | 28               |
|------------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|
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| 118              | 118              | 118              | 118              | 118              | 118              | 98               | 98               | 98               | 98               | 98               | 98               | 98               | 98               | 89               | 89               | 89               | 89               |
| 4.               | 4                | 4                | 4                | w                | ω                | 4.               | 4.               | 4.               | 4                | 4.               | 4.               | ω                | w                | 4.               | 4                | 4                | w                |
| US-10-169-003-56 | US-10-169-003-46 | US-10-182-018-56 | US-10-182-018-46 | US-09-423-800-56 | US-09-423-800-46 | US-10-733-563-39 | US-10-733-563-38 | US-10-766-610-39 | US-10-766-610-38 | US-10-766-773-39 | US-10-766-773-38 | US-09-840-499-39 | US-09-840-459-38 | US-10-733-563-49 | US-10-766-610-49 | US-10-766-773-49 | US-09-840-499-49 |
| Sequence 56,     | -                | -                | 4 1              | 56,              | 46,              |                  | -                |                  |                  |                  |                  | 9 4              |                  |                  | 49,              | 4 4              | 49,              |
| Appl             | rddy             | rddy             | Tddv             | Appl             | Appl             | Appl             | Tddv             | Iddy             | Tddv             | rddy             | Appl             |

## ALIGNMENTS

| RESULT 2 US-10-632-706-145 ; Sequence 145, Application US/10632706 ; Sequence 145, Application US/10632706 ; Publication No. US20040175385A1 ; GENERAL INFORMATION: ; APPLICANT: MARKS, JAMES D. ; APPLICANT: MARKS, JAMES D. ; APPLICANT: MARKSDORFER, PETER ; APPLICANT: MARKSDORFER, PETER ; TITLE OF INVENTION: THERAPEUTIC MONOCLONAL ANTIBODIES THAT NEUTRALIZE BOTULINUM ; TITLE OF INVENTION: NEUROTOXINS ; FILE REFERENCE: 407T-895120US ; CURRENT APPLICATION NUMBER: US/10/632,706 ; CURRENT FILING DATE: 2003-08-01 ; PRIOR APPLICATION NUMBER: US 60/400,721 ; PRIOR APPLICATION NUMBER: US 09/144,806 | Query Match 100.0%; Score 93; DB 4; Length 17; Best Local Similarity 100.0%; Pred. No. 3.3e-07; Matches 17; Conservative 0; Mismatches 0; Indels 0; Gaps 0; Qy 1 TISDGGSYTYYPDSVKG 17 |  | RESULT 1  US-10-632-706-141  US-10-632-706-141  Sequence 141, Application US/10632706  Sequence 141, Application US/10632706  Publication No. US20040175385A1  GENERAL INFORMATION:  APPLICANT: MARKS, JAMES D.  APPLICANT: AMERSDOREER, PETER  TITLE OF INVENTION: THERAPEUTIC MONOCLONAL ANTIBODIES THAT NEUTRALIZE BOTULINUM  TITLE OF INVENTION: NEUROTOXINS  TITLE OF INVENTION: NEUROTOXINS  FILE REFERENCE: 407T-895120US  CURRENT APPLICATION NUMBER: US/10/632,706  CURRENT APPLICATION NUMBER: US/10/632,706  PRIOR APPLICATION NUMBER: US/10/632,706  PRIOR APPLICATION NUMBER: US/10/632,706 |
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US-10-632-706-153
, Sequence 153, Application US/10632706
, Publication No. US20040175385A1
, GENERAL INFORMATION:
, APPLICANT: MARKS, JAMES D.
, APPLICANT: AMERSDORFER, PETER
  ; FEATURE:
; OTHER INFORMATION: single chain antibody fragment US-10-632-706-149
   ; TYPE: PRT
; ORGANISM: Artificial
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; OTHER INFORMATION: single chain antibody fragment
US-10-632-706-145
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Publication No. US20040175385A1
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NUMBER OF SEQ ID NOS: 278
SOFTWARE: PatentIn version 3.2
SEQ ID NO 145
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   GENERAL INFORMATION:
APPLICANT: AMERŚDORFER, PETER
TITLE OF INVENTION: THERAPEUTIC MONOCLONAL ANTIBODIES THAT NEUTRALIZE BOTULINUM
TITLE OF INVENTION: NEUROTOXINS
FILE REFERENCE: 407T-895120US
CURRENT FILING DATE: 2003-08-01
PRIOR APPLICATION NUMBER: US/10/632,706
CURRENT FILING DATE: 2003-08-01
PRIOR FILING DATE: 2002-08-01
PRIOR FILING DATE: 2002-08-01
PRIOR PRIOR APPLICATION NUMBER: US 60/400,721
PRIOR PRIOR APPLICATION NUMBER: US 99/144,806
PRIOR FILING DATE: 1998-08-31
NUMBER OF SEQ ID NOS: 278
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  TITLE OF INVENTION: THERAPEUTIC MONOCLONAL ANTIBODIES THAT NEUTRALIZE BOTULINUM TITLE OF INVENTION: NEUROTOXINS FILE REFERENCE: 4077-895120US CURRENT FILING DATE: 2003-08-01 PRIOR APPLICATION NUMBER: US 60/400,721 PRIOR PILING DATE: 1998-08-31 NUMBER OF SEQ ID NOS: 278
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100.0%; Pred. No.
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   Length 17;
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US-10-632-706-157
  RESULT 6
US-09-144-886-62
  US-10-632-706-157
   Sequence 62, Application US/09144886
Patent No. US20020155114A1
GENERAL INFORMATION:
APPLICANT: Marks, James D
APPLICANT: Ameridorfer, Peter
APPLICANT: Ameridorfer, Peter
TITLE OF INVENTION: Therapeutic Monoclonal
TITLE OF INVENTION: Botulinum Reurotoxins
FILE REFERENCE: 2500.117USO
  CURRENT APPLICATION NUMBER: US/09/144,886
CURRENT FILING DATE: 1998-08-31
NUMBER OF SEQ ID NOS: 98
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 62
LENGTH: 118
   SOFTWARE: PatentIn version 3.2
SEQ ID NO 157
LENGTH: 17
   Sequence 157, Applic Publication No. US20 GENERAL INFORMATION:
   SEQ ID NO 153
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Best Local Similarity
   APPLICANT: MARKS, JAMES D.
APPLICANT: AMERSDORFER, I
  OTHER INFORMATION: single chain antibody fragment
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100.0%; Pred. No. 3.3e-07;
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TISDGGSYTYYPDSVKG

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  US-10-632-706-59
  Sequence 63, Application US/09144886
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GENERAL INFORMATION:
APPLICANT: Marks, James D
APPLICANT: Mersdorfer, Peter
TITLE OF INVENTION: Therapeutic Monoclonal Antibodies That Neutralize
TITLE OF INVENTION: Botulinum Neurotoxins
FILE REFERENCE: 2500.117USO
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CURRENT APPLICATION NUMBER: US/09/144,886
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US-10-632-706-60
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Publication No. US200401753
GENERAL INFORMATION:
APPLICANT: MARKS, JAMES D.
APPLICANT: AMERSDORFER, P.
  Query Match
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Matches 17; Conserv
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   TITLE OF INVENTION: THERAPBUTIC MONOCLONAL ANTIBODIES THAT NEUTRALIZE BOTULINUM TITLE OF INVENTION: NEUROTOXINS FILE REFERENCE: 407T-895120US CURRENT PAPLICATION NUMBER: US/10/632,706 CURRENT FILING DATE: 2003-08-01 PRIOR APPLICATION NUMBER: US 60/400,721 PRIOR APPLICATION NUMBER: US 60/400,721 PRIOR APPLICATION NUMBER: US 60/400,721 PRIOR PILING DATE: 2002-08-01 PRIOR FILING DATE: 2002-08-01 PRIOR FILING DATE: 1908-08-31
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APPLICANT: MEDAREX, INC.
   ORGANISM: Artificial
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PRIOR APPLICATION DATA:
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PILING DATE: <Unknown>
ATTORNEY/AGENT INFORMATION:
   COMPUTER READABLE FORM,
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DO9
  SOFTWARE: ASCII text CURRENT APPLICATION DATA:
NAME: Arnold, Beth E.
REGISTRATION NUMBER: 35,430
REFERENCE/DOCKET NUMBER: MXI-013
TELECOMMUNICATION INFORMATION:
  NUMBER OF SEQUENCES: 28
   TITLE OF INVENTION: HUMANIZED ANTIBÓDIES TO FC RECEPTORS FOR IMMUNOBLOBULIN G ON HUMAN MONONUCLEAR PHAGOCYTES
   50 TISDGGSYTYYPDSVKG
  TISDGGSYTYYPDSVKG 17
   APPLICATION NUMBER: US/10/229,335 FILING DATE: 26-Aug-2002
   COUNTRY: USA
ZIP: 08801
  CITY: Annandale
STATE: New Jersey
   Application US/10632706
No. US20040175385A1
   STREET: P.O.
   Conservative
   100.0%; Score 93; DB 4;
100.0%; Pred. No. 2.5e-06;
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   66
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RESULT 12
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Publication No. US20030144483A1
GENERAL INFORMATION:
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SEQUENCE CHARACTERISTICS:
   INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
   APPLICATION NUMBER: US/08/435,516
FILING DATE: «Unknown»
ATTORNEY/AGENT INFORMATION:
NAME: ARTOID, Beth E.
REGISTRATION NUMBER: 35,430
REFERENCE/DOCKET NUMBER: MXI-013
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 227-7400
TELEPHONE: (617) 227-5941
  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
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SOFTWARE: ASCII text
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FILING DATE: 26-AUG-2002
CLASSIFICATION: <UNknown>
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   APPLICANT: MEDAREX, INC.
APPLICANT: MEDAREX, INC.
TITLE OF INVENTION: HUMANIZED ANTIBODIES TO FC RECEPTORS FOR
IMMUNOBLOBULIN G ON HUMAN MONONUCLEAR PHAGOCYTES
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APPLICANT: BIGGEN, INC.
APPLICANT: GARBER, E11en
APPLICANT: LYME, Paul
APPLICANT: LYME, Paul
APPLICANT: SALDHANA, JOSE W.
TITLE OF INVENTION: HUMANIZED ANTI-LT-BETA-R ANTIBODIES
FILE REFERENCE: BINA100CN
CURRENT APPLICATION NUMBER: US/10/412,406
CURRENT FILING DATE: 2003-04-10
PRIOR APPLICATION NUMBER: 60/240,285
PRIOR FILING DATE: 2000-10-13
PRIOR APPLICATION NUMBER: 60/275,289
PRIOR FILING DATE: 2001-03-13
PRIOR APPLICATION NUMBER: 60/299,987
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PRIOR FILING DATE: 2001-03-13
PRIOR FILING DATE: 2001-05-13
PRIOR FILING DATE: 2001-05-13
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PRIOR FILING DATE: 2001-10-12
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Publication No. US20030144483A1

GENERAL INFORMATION:

APPLICANT: MEDAREX, INC.

TITLE OF INVENTION: HUMANIZED ANTIBODIES TO FC RECEPTORS FOR

IMMUNOBLOBULIN G ON HUMAN MONONUCLEAR PHAGOCYTES
  Sequence 2, Application US/10412406 Publication No. US20040058394A1 GENERAL INFORMATION:
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PILING DATE: 26-Aug-2002
CLASSIFICATION: «Uhknown»
PRIOR APPLICATION NUMBER: US/08/435,516
FILING DATE: «Uhknown»
ATTORNEY/AGENT INFORMATION:
NAME: Arnold, Beth E.
REGISTRATION NUMBER: 35,430
REFERENCE/DOCKET NUMBER: MXI-013
   REFERENCE/DOCKET NUMBER:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 227-7400
   SOFTWARE: ASCII text CURRENT APPLICATION DATA:
  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
   NUMBER OF SEQUENCES: 28
STREET: P.O. Box 953,
   50 TISDGGSYTYYPDSVKG 66
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FastSEQ for Windows Version 4.0
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  TOPOLOGY: linear
   TELEFAX: (617)
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Publication No. US20040058394A1
GENERAL INFORMATION:
APPLICANT: BIOGEN, INC.
APPLICANT: SALDHAMA, JOSE W.
TITLE OF INVENTION: HUMANIZED ANTI-LT-BETA-R ANTIBODIES
FILE REFERENCE: BINALOCK
CURRENT APPLICATION NUMBER: US/10/412,406
CURRENT FILING DATE: 2003-04-10
PRIOR APPLICATION NUMBER: 60/240,285
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PRIOR REPLICATION NUMBER: 60/299,987
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PRIOR FILING DATE: 2001-06-21
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PRIOR PILING DATE: 2001-10-12
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   US-10-412-406-10
Sequence 10, Application US/10412406
Publication No. US20040058394A1
GENERAL INFORMATION:
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Best Local Similarity
Matches 17; Conserv
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Best Local Similarity 100.0%;
Matches 17; Conservative 0
  APPLICANT: LYNE, Paul
APPLICANT: SALDHANA, JOSE W.
TITLE OF INVENTION: HUMANIZED ANTI-LT-BETA-R ANTIBODIES
FILE REFERENCE: BINA100CN
CURRENT APPLICATION NUMBER: US/10/412,406
CURRENT FILING DATE: 2003-04-10
   APPLICANT: BIOGEN, INC.
APPLICANT: GARBER, Ellen
APPLICANT: LYNE, Paul
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GenCore version 5.1.6 Copyright (c) 1993 - 2005 Compugen Ltd

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and is derived by analysis of the total score distribution.
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    protein search, using sw model
   Published Applications AA New:*

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   59.1
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 US-11-172-320-6
US-11-144-248-16
US-10-789-273-1
US-10-789-273-9
US-10-789-273-1
US-10-789-273-12
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; Sequence 16, Application US/11144248
; Publication No. US20050244408A1
; GENERAL INFORMATION:
; APPLICANT: Cohen, Bruce D.
; APPLICANT: Miller, Penelope E.
; APPLICANT: Miller, James D.
; APPLICANT: Moyer, James D.

| Qy 1 TISDGGSYTYYPDSVKG 17                     Db 50 TISSGGSYTYYLDSIKG 66 | Query Match 81.7%; Score 76; DB 7; Length 444; Best Local Similarity 82.4%; Pred. No. 8.5e-06; Matches 14; Conservative 1; Mismatches 2; Indels 0; | RESULT 1 103-16 Sequence 6, Application US/11172320 Sequence 6, Application Mo. US20050244413A1 GENERAL INFORMATION: APPLICANT: Hadolf, Guenther APPLICANT: Heider, Karl-Heinz TITLE OF INVENTION: Compositions and Methods for Treating Cancer of INVENTION: Chemotherapeutic Agents TITLE OF INVENTION: Chemotherapeutic Agents FILE REFERENCE: 1/1383 CURRENT APPLICATION NUMBER: US/11/172,320 CURRENT FILING DATE: 2005-06-30 PRIOR APPLICATION NUMBER: US/10/645,215 PRIOR FILING DATE: 2003-08-21 PRIOR FILING DATE: August 21, 2002 PRIOR FILING DATE: August 21, 2002 PRIOR FILING DATE: August 21, 2002 PRIOR FILING DATE: August 21, 2002 PRIOR FILING DATE: August 26, 2002 NUMBER OF SEQ ID NOS: 9 SOFTWARE: PatentIn Ver. 2.1 SEQ ID NO 6 LENGTH: 444 TYPE: PRT ORGANISM: Artificial Sequence PEATURE: OTHER INFORMATION: Humanised Murine Antibody BIWA 4 Heavy Chain US-11-172-320-6 | ALIGNMENTS | 26 37 39.8 327 1 US-10-512-184-64 27 37 39.8 328 1 US-10-512-184-63 28 37 39.8 429 7 US-11-074-176-248 29 37 39.8 429 7 US-11-074-176-248 29 37 39.8 429 7 US-11-191-184-65 29 37 39.8 625 1 US-10-512-184-47 30 37 39.8 625 1 US-10-512-184-47 31 36 38.7 119 7 US-11-109-954-5 32 36 38.7 119 7 US-11-109-156-22 34 35 37.6 203 1 US-10-318-62-22 35 36 38.7 142 7 US-11-199-156-22 36 37.6 203 1 US-10-310-386-122 37 39.8 203 1 US-10-131-826A-520 38 37.6 218 7 US-11-190-9643-4 39 34.5 37.1 133 7 US-11-010-748A-3 40 34.5 37.1 133 7 US-11-010-748A-3 41 34.5 37.1 133 7 US-11-010-748A-3 42 34.5 37.1 133 7 US-11-010-748A-3 43 34.6 313 1 US-10-512-184-35 36 31.6 31.6 1 US-10-512-184-35 37 36.6 31.1 1 US-10-982-545-11 38 39.8 267 US-10-982-545-11 39 39 39 39 39 39 39 39 39 39 39 39 39 3                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              |
|--------------------------------------------------------------------------|----------------------------------------------------------------------------------------------------------------------------------------------------|------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|------------|-----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
|                                                                          | .,                                                                                                                                                 | cer                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  |            | sequence sequence sequence sequence sequence sequence sequence sequence sequence sequence sequence sequence sequence sequence                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           |
|                                                                          | Gaps                                                                                                                                               | g using                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              |            | 64, Appl<br>63, Appl<br>64, Appl<br>65, Appl<br>47, Appl<br>462, Appl<br>462, Appl<br>27, Appl<br>27, Appl<br>28, Appl<br>29, Appl<br>21, Appl<br>31, Appl<br>41, Appl<br>4 |
|                                                                          | 0;                                                                                                                                                 |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      |            |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         |

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FILE REFERENCE: ELN-002CP
CURRENT APPLICATION NUMBER: US/10/789,273
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PRIOR APPLICATION NUMBER: US/10/388,389
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PRIOR FILING DATE: 2001-12-06
PRIOR FILING DATE: 2001-12-06
PRIOR APPLICATION NUMBER: US 60/251,892
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PRIOR PILING DATE: 2000-12-06
NUMBER OF SEQ ID NOS: 63
SOFTMARE: FastSEQ for Windows Version 4.0
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Publication No. US20050244408A1
GENERAL INFORMATION:
APPLICANT: Cohen, Bruce D.
APPLICANT: Beebe, Jean
APPLICANT: Miller, Penelope E.
APPLICANT: Moyer, James D.
   PRIOR FILING DATE: 2001-01-05
NUMBER OF SEQ ID NOS: 60
SOFTWARE: PatentIN Ver. 2.1
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APPLICANT: Basi, Guriq
APPLICANT: Saldanha, Jose
APPLICANT: Yednock, Ted
TITLE OF INVENTION: HUMANIZED ANTIBODIES THAT RECOGNIZE
TITLE OF INVENTION: BETA-AMYLOID PEPTIDE
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APPLICANT: Saldanha, Jose
APPLICANT: Saldanha, Jose
APPLICANT: Saldanha, Jose
APPLICANT: Saldanha, Ted
TITLE OF INVENTION: HUMANIZED ANTIBODIES THAT RECOGNIZE
TITLE OF INVENTION: BETA-AMYLOID PRFTIDE
FILE REFERENCE: ELN-02CP
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CURRENT FILING DATE: 2004-02-27
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US-10-789-273-9
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APPLICANT: Bas1, Guriq
APPLICANT: Saldanha, Jose
APPLICANT: Vednock, Tose
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TITLE OF INVENTION: HUMANIZED ANTIBODIES THAT RECOGNIZE
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CURRENT APPLICATION NUMBER: US/10/789,273
CURRENT FILING DATE: 2004-02-27
PRIOR APPLICATION NUMBER: US/10/388,389
PRIOR FILING DATE: 2003-03-12
PRIOR FILING DATE: 2003-03-12
PRIOR FILING DATE: 2001-12-06
PRIOR APPLICATION NUMBER: US 60/251,892
PRIOR APPLICATION NUMBER: US 60/251,892
PRIOR FILING DATE: 2000-12-06
NUMBER OF SEQ ID NOS: 63
SOFTWARE: FastESEQ for Windows Version 4.0
SEQ ID NO 4
  APPLICANT: Basi, Guriq
APPLICANT: Saldanha, Jose
APPLICANT: Saldanha, Jose
APPLICANT: Saldanha, Jose
APPLICANT: Saldanha, Jose
TITLS OF INVENTION: HUMANIZED ANTIBODIES THAT RECOGNIZE
TITLE OF INVENTION: BETA-AMYLOID PEPTIDE
FILE REFERENCE: ELN-OO2CP
CURRENT APPLICATION NUMBER: US/10/789,273
CURRENT FILING DATE: 2004-02-27
PRIOR APPLICATION NUMBER: US/10/388,389
PRIOR APPLICATION NUMBER: US/10/388,389
PRIOR APPLICATION NUMBER: US/10/388,389
PRIOR APPLICATION NUMBER: US/10/388,389
PRIOR FILING DATE: 2003-03-12
PRIOR FILING DATE: 2003-03-12
PRIOR FILING DATE: 2003-03-12
PRIOR FILING DATE: 2001-12-06
RESULT 8
US-10-789-273-12
; Sequence 12, Application US/10789273
; Publication No. US20050249725A1
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   US-10-789-273-4
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Best Local Similarity be.
  Query Match 58.1
Best Local Similarity 58.8
Matches 10; Conservative
   PRIOR APPLICATION NUMBER: US 60/251,892
PRIOR FILING DATE: 2000-12-06
NUMBER OF SEQ ID NOS: 63
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SEQ ID NO 8
  Publication No.
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LOCATION: (1)...(19)
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FEATURE:
OTHER INFORMATION: Humanized 3D6 heavy chain variable region
FEATURE:
NAME/KEY: SIGNAL
LOCATION: (1)...(19)
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ORGANISM: Mus musculus
   TYPE: PRT ORGANISM: Artificial Sequence
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   1 TISDGGSYTYYPDSVKG 17
  Application US/10789273
No. US20050249725A1
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Pred. No.
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  DB 1; Length 138; 0.0075;
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  0;
   Gaps
  Gaps
   0
   0
```

```
APPLICANT: Saldanha, Jose
APPLICANT: Saldanha, Jose
APPLICANT: Saldanha, Jose
APPLICANT: Yednock, Ted
APPLICANT: Yednock, Ted
TITLE OF INVENTION: BETA-AMYLOID PEPTIDE
FILE REFERENCE: ELN-002CP
CURRENT APPLICATION NUMBER: US/10/789,273
CURRENT APPLICATION NUMBER: US/10/388,389
PRIOR APPLICATION NUMBER: US 10/010,942
PRIOR APPLICATION NUMBER: US 10/010,942
PRIOR APPLICATION NUMBER: US 60/251,892
PRIOR FILING DATE: 2001-12-06
PRIOR PILING DATE: 2001-12-06
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ORGANISM: Artificial Sequence
FEATUURE:
OTHER INFORMATION: Humanized 3D6 light chain variable reg
  ; FEATURE:
; NAME/KEY: SIGNAL
; LOCATION: (1)...(19)
US-10-789-273-12
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  RESULT 9
US-11-144-248-46
   US-11-144-248-46
  GENERAL INFORMATION:
  Sequence 46, Appropriation No.
  GENERAL INFORMATION:
  Matches
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  Query Match
Best Local S
   SEQ ID NO 46
  APPLICANT: Miller, Penelope E.
APPLICANT: Moyer, James D.
APPLICANT: Moyer, James D.
APPLICANT: Corvellan, Jose R.
APPLICANT: Gallo, Michael
TITLE OF INVENTION: ANTIBODIES TO INSULIN-LIKE GROWTH FACTOR I RECEPTOR
FILE REFERENCE: ABX-PF2
CURRENT APPLICATION NUMBER: US/11/144,248
CURRENT FILING DATE: 2005-06-02
PRIOR APPLICATION NUMBER: US/10/038,591
PRIOR APPLICATION NUMBER: US/10/038,591
PRIOR PILING DATE: 2002-01-04
   PRIOR APPLICATION NUMBER: 60/259,927
PRIOR FILING DATE: 2001-01-05
NUMBER OF SEQ ID NOS: 60
  APPLICANT:
  APPLICANT: Cohen, Bruce D.
   SOFTWARE: PatentIn Ver. 2.1
  LENGTH: 470
TYPE: PRT
ORGANISM: Homo sapiens
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  Match 58.1%;
Local Similarity 58.8%;
see 10; Conservative
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   1 TISDGGSYTYYPDSVKG 17
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  ISGSGGSTYYADSVKG
                                     ISDGGSYTYYPDSVKG 17
  Application US/11144248
  US20050244408A1
  85
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; ORGANISM: Homo sapiens
US-11-144-248-30
   US-11-144-248-50
  SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 50
LENGTH: 473
   Sequence 50, Application US/11144248 Publication No. US20050244408A1 GENERAL INFORMATION:
   SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 30
   GENERAL INFORMATION:
APPLICANT: Cohen, Bruce D.
APPLICANT: Beebe, Jean
   Sequence 30, Application US/11144248 Publication No. US20050244408A1
   Query Match
   Query Match
Best Local Similarity
   CURRENT APPLICATION NUMBER: US/11/144,248
CURRENT FILING DATE: 2005-06-02
PRIOR APPLICATION NUMBER: US/10/038,591
PRIOR FILING DATE: 2002-01-04
PRIOR PELICATION NUMBER: 60/259,927
PRIOR FILING DATE: 2001-0-05
PRIOR FILING DATE: 2001-0-05
NUMBER OF SEQ ID NOS: 60
  CURRENT APPLICATION NUMBER: US/11/144,248
CURRENT FILING DATE: 2005-06-02
PRIOR APPLICATION NUMBER: US/10/038,591
PRIOR FILING DATE: 2002-01-04
   APPLICANT: Beebe, Jean
APPLICANT: Miller, Penelope E.
APPLICANT: Moyer, James D.
APPLICANT: Corvalan, Jose R.
APPLICANT: Gallo, Michael
APPLICANT: Gallo, Michael
TITLE OF INVENTION: ANTIBODIES TO INSULIN-LIKE GROWTH FACTOR I RECEPTOR
   APPLICANT: Cohen, Bruce D. APPLICANT: Beebe, Jean APPLICANT: Miller, Penelog
   PRIOR APPLICATION NUMBER: 60/259,927
PRIOR FILING DATE: 2001-01-05
NUMBER OF SEQ ID NOS: 60
  APPLICANT: COTVAIAN, JOSE R.
APPLICANT: Gallo, Michael
TITLE OF INVENTION: ANTIBODIES TO INSULIN-LIKE GROWTH FACTOR I RECEPTOR
FILE REFERENCE: ABX-PF2
   APPLICANT: Miller, Penelope E. APPLICANT: Moyer, James D.
  11-144-248-30
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  70
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Conservative
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Pred. No. 0.04;
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RESULT 12

RESULT 14 US-10-512-184-27

```
APPLICANT: Heebs, Jean
APPLICANT: Miller, Penelope E.
APPLICANT: Moyer, James D.
APPLICANT: Corvalan, Jose R.
APPLICANT: Gorian, Jose R.
APPLICANT: Gallo, Michael
TITLE OF INVENTION: ANTIBODIES TO INSULIN-LIKE GROWTH FACTOR I RECEPTOR
FILE REFERENCE: ABX-PP2
CURRENT APPLICATION NUMBER: US/11/144,248
CURRENT FILING DATE: 2005-06-02
PRIOR APPLICATION NUMBER: US/10/038,591
PRIOR APPLICATION NUMBER: 60/259,927
PRIOR FILING DATE: 2001-01-04
PRIOR FILING DATE: 2001-01-05
NUMBER: OF SEQ. ID NOS: 60
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밁
   US-11-144-248-24
   US-11-144-248-24
  ; ORGANISM: Homo sapiens
US-11-144-248-8
  US-11-144-248-8
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TYPE: PRT
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  Sequence 8, Application US/11144248
Publication No. US20050244408A1
GENERAL INFORMATION:
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Publication No. US20050244408A1
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CURRENT APPLICATION NUMBER: US/11/144,248
CURRENT FILING DATE: 2005-06-02
PRIOR APPLICATION NUMBER: US/10/038,591
PRIOR FILING DATE: 2002-01-04
PRIOR FILING DATE: 2002-01-04
PRIOR PPLICATION NUMBER: 60/259,927
PRIOR APPLICATION NUMBER: 60/259,927
PRIOR FILING DATE: 2001-01-05
NUMBER OF SEQ ID NOS: 60
   SOFTWARE: Patentin Ver. 2.1
  SOFTWARE: PatentIn Ver. 2.1
  APPLICANT:
   APPLICANT: Cohen, Bruce D. APPLICANT: Beebe, Jean
  APPLICANT: Cohen, Bruce D.
   APPLICANT:
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   INFORMATION:
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50 ISGSGGTTFYADSVKG 65
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: Moyer, James D.
: Corvalan, Jose R.
: Gallo, Michael
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Pred. No.
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Publication No. US20050244901A1

Publication No. US20050244901A1

GENERAL INFORMATION:

APPLICANT: Fraunhofer Gesellschaft zur F"rderung der angewandten Forschung e.V.

TITLE OF INVENTION: Antibodies, recombinant antibodies, recombinant

TITLE OF INVENTION: Antibody fragments and fusions mediated plant disease

TITLE OF INVENTION: resistance against fungi

PILE REFERENCE: 3581.01US01

CURRENT APPLICATION NUMBER: US/10/512,184

CURRENT FILING DATE: 2004-10-22

NUMBER OF SEQ ID NOS: 72

SOFTWARE: Patentin Ver. 2.1

SEQ ID NO 27

LENGTH: 250

TYPE: PRT
ORGANISM: Artificial Sequence
PEATURE: OFTHER INFORMATION: Description of Artificial Sequence: scFv SGB3 with
OTHER INFORMATION: specificity against Fusarium ssp.; oxiginates from
OTHER INFORMATION: Gallus gallus.

US-10-512-184-27
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Publication No. US20050244408A1
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APPLICANT: Cohen, Bruce D.
APPLICANT: Beabe, Jean
APPLICANT: Miller, Penelope E.
APPLICANT: Myer, James D.
   RESULT 15
US-11-144-248-45
   밁
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-144-248-45
Search completed: November 21, 2005, 12:33:52
Job time : 1.41136 secs
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CURRENT PILING DATE: 2005-06-02
PRIOR APPLICATION NUMBER: US/11/144,248
CURRENT FILING DATE: 2002-01-04
PRIOR FILING DATE: 2002-01-04
PRIOR FILING DATE: 2002-01-04
PRIOR APPLICATION NUMBER: 60/259,927
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   Corvalan, Jose R.
Gallo, Michael
   85
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  geneseqp2005s:*
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| 45             | 44             | 43       | 42       | 41       | 40       | 39       | 38       | 37       | 36       | 3<br>5   | 34       | y<br>y   | 32       | 31       | 30       | 29       | 28       | 27       | 26       | 25       |
|----------------|----------------|----------|----------|----------|----------|----------|----------|----------|----------|----------|----------|----------|----------|----------|----------|----------|----------|----------|----------|----------|
| 138            | 138            | 138      | 138      | 138      | 138      | 138      | 138      | 138      | 138      | 138      | 138      | 138      | 138      | 138      | 138      | 138      | 138      | 138      | 138      | 138      |
| 89.6           | 89.6           | 89.6     | 89.6     | 89.6     | 89.6     | 89.6     | 89.6     | 89.6     | 89.6     | 89.6     | 89.6     | 89.6     | 89.6     | 89.6     | 89.6     |          |          |          | 89.6     |          |
| 115            | 115            | 115      | 115      | 115      | 115      | 115      | 115      | 115      | 115      | 115      | 115      | 115      | 115      | 115      | 115      | 115      | 115      | 115      | 115      | 115      |
| 9              | 9              | 9        | 9        | 9        | ø        | 9        | 9        | 9        | 9        | φ        | 9        | 9        | 9        | 9        | ø        | ø        | 9        | 9        | 9        | 9        |
| AEA08986       | AEA08984       | AEA08982 | AEA08901 | AEA08899 | AEA08985 | AEA08981 | AEA08897 | AEA08990 | AEA08915 | AEA08987 | AEA08895 | AEA08995 | AEA08903 | AEA08989 | AEA08994 | AEA08983 | AEA08913 | AEA08911 | AEA08988 | ABA08905 |
| деа08986 неачу | Aea08984 Heavy |          | -        | _        |          |          |          |          | -        |          |          |          |          |          |          |          |          |          |          |          |
|                | -              | _        | •        | -        | •        |          |          |          |          |          |          |          |          | •        | -        | -        | - 1      |          | cna      | cha      |

# ALIGNMENTS

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  31-AUG-1998; 98US-00144886.
01-AUG-2002; 2002US-0400721P.
   Mus
   antibacterial; antibody; botulinum neurotoxin type A; BONT/A;
BONT/A neutralising epitope; anti-botulinum neurotoxin antibody;
toxin neutralisation; botulinum neurotoxin poisoning; mouse; heavy chain;
  ADR38764 standard; peptide; 30 AA.
  01-AUG-2003; 2003US-00632706.
  09-SEP-2004.
   US2004175385-A1
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   Mouse heavy
  02-DEC-2004
  ADR38764;
   ф
   chain anti-BoNT-antibody framework 1 seqid
  (first entry)
  166
```

The invention describes an isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a specific clone where (I) binds to and neutralises botulinum neurotoxin type A (BoNT/A). An isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a clone chosen from clone \$25, C25, C39, IC6, 3D12, B4, IF3, huC25, A1, A22, WR1(V), WR1(T), 3-1, 3-8, 3-10 and ING1, where (I) binds to and neutralizes botulinum neurotoxin type A (BoNT/A). Also described are: a polypeptide (II) comprising BoNT/A neutralising epitope having an epitope that is

New isolated antibody that neutralizes botulinum neurotoxin type A, useful for diagnosing botulism or for treating pathologies associated with botulinum neurotoxin poisoning.

WPI; 2004-652009/63.

Marks JD,

Amersdorfer

P,

(REGC ) UNIV CALIFORNIA.

Example 4; SEQ ID NO 166; 110pp; English.

```
5
  8888888888888888888888
  밁
   specifically bound by an antibody expressed by clones as mentioned in (I) producing (I); and a composition (III) comprising several anti-
controllinum neurotoxin antibodies, where each antibody is specific for a different epitope of a botulinum neurotoxin and the combination of a ntibodies shows greater toxin neutralisation than the single antibodies in surplus. The following are disclosed: a pharmaceutical composition comprising (I); and a kit comprising (I); is useful for neutralising bont/A antibody and for neutralising a botulinum neurotoxin which involves contacting neurotoxin with (I) in surplus, where each of (I) is specific for a different epitope of the botulinum neurotoxin and the combination of antibodies shows greater toxin neutralisation than the single antibodies in surplus. (I) is useful for diagnosing the botulism or for treating pathologies associated with botulinum neurotoxin pathologies associated with botulinum neurotoxin consing. (I) exhibits specificity and affinity towards BoNT/A. (I) enables rapid detection or diagnosis of botulism. This is the amino acid sequence of mouse heavy chain anti-BoNT-antibody framework region 1.
   Query Match
Best Local S
Matches 30
  New isolated antibody that neutralizes botulinum neurotoxin type A, useful for diagnosing botulism or for treating pathologies associated
   antibacterial; antibody; botulinum neurotoxin type A; BoNT/A;
BoNT/A neutralising epitope; anti-botulinum neurotoxin antibody;
toxin neutralisation; botulinum neurotoxin poisoning; mouse; heavy chain;
   WPI; 2004-652009/63.
   Marks
  31-AUG-1998; 98US-00144886.
01-AUG-2002; 2002US-0400721P.
   01-AUG-2003; 2003US-00632706.
  09-SEP-2004.
   US2004175385-A1
  framework 1.
   Mouse heavy chain anti-BoNT-antibody framework 1 seqid 158
   02-DEC-2004
  ADR38756;
  ADR38756 standard; peptide;
  Sequence 30 AA;
  (REGC ) UNIV CALIFORNIA.
   Ą
   30;
  1 QVQLVQSGGGVVHPGRSLKLSCAGSGFTFS 30
   Similarity
   Amersdorfer
   (first entry)
   Conservative
   100.0%;
   30
   0;
   Score 154; DB 8;
Pred. No. 2.5e-13;
   Mismatches
  S
O
   Length 30
   Indels
   ,
,
   Gaps
```

The invention describes an isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a specific clone where (I) binds to and neutralises botulinum neurotoxin type A (BONT/A). An isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a clone chosen from clone \$25, C25, C39, IC6, 3D12, B4, IF3, huC25, Ar1, Ar2, WR1(V), WR1(T), 3-1, 3-8, 3-10 and ING1, where (I) binds to and neutralizes botulinum

The invention describes an isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a specific clone where (I) binds to and neutralises botulinum neutrotoxin type A (BoNT/A). An isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a clone chosen from clone

Example

4

SEQ ID NO 170; 110pp; English

Example 4;

SEQ ID NO 158; 110pp; English.

botulinum

neurotoxin

poisoning.

```
ADRIBUT 3
ADRIBUT 3
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   S
   comprising BoNT/A neutralising epitope having an epitope that is specifically bound by an antibody expressed by clones as mentioned in (I) producing (I); and a composition (II) comprising several anti-
comprising several anti-
composition (III) comprising several anti-
composition (III) comprising several anti-
composition (III) comprising several anti-
composition each antibodies, where each antibody is specific for a composition of the shows greater toxin neutralisation than the single antibodies shows greater toxin neutralisation than the single antibodies in surplus. The following are disclosed: a pharmaceutical composition comprising (I); and a kit comprising (I). (I) is useful for neutralising COMPIA antibody and for neutralising a botulinum neutrotoxin which involves contacting neurotoxin with (I) in surplus, where each of (I) is specific for a different epitope of the botulinum neutralisation than the complication of antibodies shows greater toxin neutralisation than the single antibodies in surplus. (I) is useful for diagnosing the botulism correction or for treating pathologies associated with botulinum neurotoxin correction of antibodies and affinity towards BoNT/A. (I) enables rapid detection or diagnosis of botulism. This is the amino acid sequence of mouse heavy chain anti-BoNT-antibody framework region 1.
  Query Match
Best Local S
Matches 30
   New isolated antibody that neutralizes botulinum neurotoxin type A, useful for diagnosing botulism or for treating pathologies associated with botulinum neurotoxin poisoning.
   WPI;
   31-AUG-1998; 98US-00144886.
01-AUG-2002; 2002US-0400721P.
  01-AUG-2003; 2003US-00632706
   antibacterial; antibody; botulinum neurotoxin type A; BoNT,
BoNT/A neutralising epitope; anti-botulinum neurotoxin ant;
   Sequence 30
   Marks
   09-SEP-2004.
   US2004175385-A1
   Mus sp
   Mouse heavy
   02-DEC-2004
  ADR38768;
  ADR38768 standard;
   (REGC
  toxin neutralisation; botulinum
  á,
   ) UNIV CALIFORNIA
   30;
  1 QVQLVQSGGGVVHPGRSLKLSCAGSGFTFS
   Similarity
   QVQLVQSGGGVVHPGRSLKLSCAGSGFTFS
  Amersdorfer
   AA,
   chain anti-BoNT-antibody framework 1 seqid 170.
   Conservative
  (first entry)
   peptide;
  100.0%;
  30
  0;
  Score 154; DB 8;
Pred. No. 2.5e-13;
Mismatches 0;
  ₹
   neurotoxin poisoning; mouse; heavy chain;
   30
  Length
   antibody;
  0
  Gaps
  0
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**អនុសម្ភាស់ អនុសម្ភាស់ អនុសម្ភាស់ អនុសម្ភាស់ អនុ** 

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RESULT 4
ADR38706
ID ADR3870
XX ADR337
XX ADR37
XX ADR37
XX MOUS
XX MOUS
XX HOUS
XX HOUS
XX US20
XX US20
XX O9-S
XX O9-S
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  CC 325, C25, C39, 1C6, 3D12, B4, 1F3, huC25, Ar1, Ar2, WR1(V), WR1(T), 3-1, CC 3-8, 3-10 and ING1, where (I) binds to and neutralizes botulinum CC neurotoxin type A (BoNT/A). Also described are: a polypeptide (II) CC comprising BoNT/A neutralising epitope having an epitope that is CC specifically bound by an antibody expressed by clones as mentioned in (I) producing (I), and a composition (II) comprising several anticoccomposition (II) comprising several anticoccomposition (II) comprising several anticoccomprising (I), and a composition and the combination of CC in surplus. The following are disclosed: a pharmaceutical composition CC comprising (I); and a kit comprising (I) is useful for neutralising CC involves contacting neurotoxin with (I) in surplus, where each of (I) is specific for a different epitope of the botulinum neurotoxin and the CC single antibodies in surplus. (I) is useful for diagnosing the botulism CC poisoning. (I) exhibits specificity and affinity towards BoNT/A. (I) canables rapid detection or diagnosis of botulism. This is the amino acid sequence of mouse heavy chain anti-BoNT-antibody framework region 1.
   Matches
  Query Match
Best Local (
  Mue
   Sequence 30 AA;
   antibacterial; antibody; botulinum neurotoxin type A; BoNT/A; BoNT/A neutralising epitope; anti-botulinum neurotoxin antibody; toxin neutralisation; botulinum neurotoxin poisoning; mouse; heavy chain;
  ADR38760 standard; peptide; 30
  31-AUG-1998;
01-AUG-2002;
   01-AUG-2003; 2003US-00632706.
  09-SEP-2004.
   Mouse heavy chain anti-BoNT-antibody framework 1 segid 162.
  02-DEC-2004 (first entry)
  New isolated antibody that neutralizes botulinum neurotoxin type A, useful for diagnosing botulism or for treating pathologies associated
  WPI; 2004-652009/63.
   US2004175385-A1
   (REGC ) UNIV CALIFORNIA.
   framework 1.
  Local Similarity
   ch 100.0%;
1 Similarity 100.0%;
30; Conservative 0
  ۳
  QVQLVQSGGGVVHPGRSLKLSCAGSGFTFS 30
  QVQLVQSGGGVVHPGRSLKLSCAGSGFTFS 30
   Amersdorfer P;
  98US-00144886.
2002US-0400721P.
   0
   Score 154; DB 8;
Pred. No. 2.5e-13;
Mismatches 0
  ⋛
   0
  Length 30;
   Indels
   0,
   Gaps
   0;
```

The invention describes an isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a specific clone where (I) binds to and neutralises botulinum neurotoxin type A

New human antibody specific for human interleukin-12 (IL-12) used to treat disorders characterized by aberrant IL-12 expression e.g. Crohn's

Example 4; SEQ ID NO 162; 110pp; English

botulinum neurotoxin poisoning.

문 Ś

```
RESULT 5
AAB40091
   CC (BONT/A). An isolated antibody (I) that specifically binds to an epitope CC specifically bound by an antibody expressed by a clone chosen from clone CS 25, C25, C39, IC6, 3D12, B4, IF3, huC25, Ar1, Ar2, WR1(V), WR1(T), 3-1, CC 3-8, 3-10 and ING1, where (I) binds to and neutralizes botulinum CC enurctoxin type A (BONT/A). Also described are: a polypeptide (II) CC comprising BONT/A neutralising epitope having an epitope that is CC specifically bound by an antibody expressed by clones as mentioned in (I); producing (I); and a composition (III) comprising several antical continuum neurotoxin antibodies, where each antibody is specific for a CC different epitope of a botulinum neurotoxin and the combination of CC comprising (I); and a kit comprising (I) is useful for neutralising CC insurplus. The following are disclosed: a pharmaceutical composition CC comprising (I); and a kit comprising (I). (I) is useful for neutralising BONT/A antibody and for neutralising a botulinum neurotoxin which CC involves contacting neurotoxin with (I) in surplus, where each of (I) is specific for a different epitope of the botulinum neurotoxin and the CC single antibodies in surplus. (I) is useful for diagnosing the botulism or for treating pathologies associated with botulinum neurotoxin than the composition or for treating pathologies associated with botulinum neurotoxin (I) exhibits specificity and affinity towards BONT/A. (I) canables rapid detection or diagnosis of botulism. This is the amino acid sequence of mouse heavy chain anti-BONT-antibody framework region 1.
밁
  8
   Query Match
Best Local S
Matches 30
   Sequence 30
   multiple sclerosis;
   complementarity determining region; CDR; antirheumatic; antiarthritic; antisclerotic; neuroprotective; antipsoriatic; antiasthmatic; cardiant; antipsarastic; antibacterial; immunosuppressive; Crohn's disease; multiple sclerosis; rheumatoid arthritis.
  Human; neutralising antibody; interleukin-12; complementarity determining region; CDR; anti-
   Anti-hIL12 antibody H chain V region amino acid sequence SEQ ID
  AAB40091 standard; protein; 97
   WPI; 2000-638250/61.
  Kaymakcalan Z, Labkovsky
Veldman GM, Venturini A,
Derbyshire EJ, Carmen S,
  Salfeld JG,
  25-MAR-1999;
   24-MAR-2000; 2000WO-US007946.
  28-SEP-2000
   WO200056772-A1
  Homo sapiens.
   05-FEB-2001
  AAB40091;
   (BADI ) BASF AG. (GEMY ) GENETICS INST INC.
   30;
   Similarity
   QVQLVQSGGGVVHPGRSLKLSCAGSGFTFS 30
   QVQLVQSGGGVVHPGRSLKLSCAGSGFTFS 30
   Ŗ,
  Z, Labkovsky B, Sakorafas P, Friec
Venturini A, Warne NW, Widom A, I
U, Carmen S, Smith S, Holtet TL,
   (first entry)
  99US-0126603P
   100.0%; Score 154; DB 8; 100.0%; Pred. No. 2.5e-13; ... Mismatches 0;
  jee S, Tracey DB, White M
Friedrich S, Myles A;
m A, Elvin JG, Duncan AR;
t TL, Du Fou SL;
   IL-12; antiinflammatory;
  Length 30;
   Gaps
  617.
   0
```

Claim 75; Page 121; 377pp; English.

WPI; 2005-252952/26.

```
RESULT 6
ADY93868
ID 93868
ID 93868
ADY93868
AC ADY9
AC ADY9
AC ADY9
AC ADY9
AC ANTI
AC ANTI
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   5
   Matches
   Query Match
Best Local &
  This invention relates to a new human antibody specific for human interlaukin-12 (II-12). The invention also includes antigen binding portions that bind to II-12. Sequences AAB39485-B39516 represent human anti-IL-12 antibody heavy and light chain complementarity determining region (CDR) amino acid sequences, and also includes variable region amino acid sequences. Other variable region amino acid sequences. Other variable region amino acid sequences and AB39517-B39560 and AAB40068-B40149. Sequences AAB39577-B40063 represent anti-II-12 CDR3 related amino acid sequences, AAB39772-B40063 represent other CDR3 sequences. Light chain CDR3 consensus sequences are given in AAB40064-B40067. Primers used in the identification and construction of the antibodies of the invention are given in AAC61062-C61071. The antibody of the invention is a neutralising antibody and has
  antirheumatic; antiarthritic; antisclerotic; antiinflammatory; neuroprotective; antipsoriatic; antiasthmatic; cardiant; antiparasitic; antibacterial and immunosuppressive activity. The antibodies or antigenbinding fragments are useful in the treatment of disorders associated with detrimental release of human IL-12, especially Crohn's disease, multiple sclerosis and rheumatoid arthritis. They can also be used in the manufacture of a pharmaceutical composition to treat human IL-12
  04-AUG-2003; 2003US-0492529P.
09-0CT-2003; 2003US-051025L9.
18-FEB-2004; 2004US-0545670P.
26-APR-2004; 2004US-0565595P.
   ADY93868 standard; protein; 97
   Sequence 97 AA;
  Ambrosino D,
Somasundaran
  (LUZU/)
   (AMBR/)
(HERN/)
(GREE/)
   04-AUG-2004; 2004US-00911838.
  US2005069869-A1
   SARS coronavirus
   glycoprotein; respiratory-gen.; virucide; sars
antibody; heavy chain variable region.
  Anti-SARS DP44 germline antibody VH region BQ ID NO:18.
  16-JUN-2005 (first entry)
  ADY93868;
   /) AMBROSINO D.
/) HERNANDEZ H.
/) GRESNOUGH T.
/) LUZURIAGA K.
/) SOMASUNDARAN F.
/) BABCOCK G J.
/) THOMAS W D.
/) SULLIVAN J.
  -
   1 QVQLVQSGGGVVHPGRSLKLSCAGSGFTFS 30
   26; Conservative
  Similarity
  EVQLVQSGGGLVHPGGSLRLSCAGSGFTFS 30
  Z,
   Hernandez H,
M, Babcock GJ,
   89.6%;
   Greenough T,
I, Thomas WD,
  Score 138; DB 3;
Pred. No. 1.1e-10;
3; Mismatches 1
   ۶
  Luzuriaga K;
Sullivan J;
  coronavirus infection;
   Length 97;
   0
   0
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Query Match Best Local S Matches 26

Similarity

89.6%;

Score 138; DB 9; Pred. No. 1.1e-10; Mismatches

Length 97;

<u>.</u> Gaps

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Sequence 97

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1 EVQLVQSGGGLVHPGGSLRLSCAGSGFTFS 30

QVQLVQSGGGVVHPGRSLKLSCAGSGFTFS 30

```
The invention relates to an isolated nucleic acid comprising a sequence concoding a SARS coronavirus (SARS-CoV) spike injectoprotein (Sprotein), or the sequence has been optimized for expression in a comprising the above nucleic acid; (2) an isolated cell comprising the comprising the above nucleic acid; (2) an isolated cell comprising the expression vector of (1); (3) a polypeptide encoded by the above nucleic cetter (2); (3) a polypeptide encoded by the above nucleic cetter (4) an isolated polypeptide comprising an extracellular portion of cetter (4) an isolated polypeptide comprising an extracellular portion of cetter (4) an isolated antibody, or its antigen inding fragment, that specifically binds to the polypeptide of (3); (6) cetter (5); (10) an isolated antibody, or its antigen binding fragment, that specifically binds to an antibody, or its antigen binding fragment, and a pharmaceutical carrier; (8) making an Scattigen binding fragment, that specifically binds to an Sprotein of SARS-CoV; (10) evaluating an antibody, or its antigen binding fragment, that specifically binds to an Sprotein of SARS-COV; (11) evaluating an antibody, or its antigen binding fragment, of SARS-COV; (12) treating a subject at risk for, (2) evaluating an antibody, or its antigen binding fragment, for comprising the antibody, or its antigen binding fragment, for comprising the antibody, or its antigen binding fragment, for comprising the antibody, or its antigen binding fragment, for comprising the antibody of (5); (13) an isolated nucleic acid confolecule encoding the antibody, its antigen-binding portion, variable the expression vector (14); (16) a host cell comprising the antibody of (5); (18) an spot cell comprising human comprising the antibody of (5); (18) an host cell comprising human communication and or vector (14); (15) an expression vector (15); (17) a transgenic mammal comprising human comprising the mammal of (14); (16) a host cell comprising an expression vector (15); (17) a transgenic mammal comprising human comprisi
   New isolated nucleic acid encoding a SARS-COV spike glycoprotein polypeptide, and optimized for expression in a human host, useful for diagnosing, preventing and/or treating SARS-COV infection.
   Example 17; SEQ ID NO 18; 138pp; English.
```

```
PRINCE EX NO. SERVING A SE
   HIV infection; anti-hiv; infection; inflammation; antiinflammatory; cancer; cytostatic; antibody; chimeric antibody; heavy chain variable region; antibody engineering; anti-IL-4R.
   Human anti-IL-4R heavy chain variable region,
  WO2005047325-A2
  28-JUL-2005
   AEA11024 standard; protein; 114 AA
  Homo sapiens.
  AEA11024;
  (first entry)
  SEQ ID NO:66
```

04-NOV-2004; 2004WO-US037241.

```
RESULT 8
ABB0725
ID ABB0
XX ABB0
XX ABB0
XX ABB0
XX ABB0
XX ABB0
XX Anti
XX Anti
XX Huma
XW Anti
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  The invention relates to an isolated polypeptide (I) comprising (a) one CC ARA10972, and antibody heavy chain variable region, or (b) ARA10978, and antibody heavy chain variable region. In (I), the antibody heavy chain or CC light chain variable region is a cynomolgus monkey antibody heavy chain or CC variable region or cynomolgus monkey antibody heavy chain variable region is a human or mouse antibody heavy chain variable region or light chain variable region. The antibody heavy chain variable region or light chain variable region is a human or mouse antibody heavy chain variable region or light chain variable comprising a sequence encoding (I); an isolated antibody (A1) comprising CC afirst polypeptide having SEQ ID No. 6, 8, 10, 12, 14 or 20, and a comprising of an antibody comprising introducing into a cynomolgus monkey a chimeric antibody and light chain and heavy chain variable regions from a cynomolgus monkey, and evaluating the effects of the cynomolgus monkey, and evaluating the effects of the chimeric antibody in the cynomolgus monkey, and evaluating the effects of the chimeric antibody in the cynomolgus monkey, and evaluating the effects of the chimeric antibody in the cynomolgus monkey, and expression vector (V1) comprising (I1); a cell (III) comprising V1; and a method of producing (I). (A1) further comprises a human antibody heavy chain variable region and a human constitute antibody light chain variable region and a human constitute antibody light chain variable region and a human constitute antibody heavy chain variable region and a human constitute antibody heavy chain variable region and a human constitute antibody. (II) (A1) further comprising V1; and a method of producing (I). (A1) further comprising V1 and a method of producing (I) (A1) further comprising V1 and a method of producing (I) (A1) further comprising V1 and a method of producing (I) (A1) further comprising V1 and a method of producing (I) (A1) further constitute (III) (III) (III) (III) (III) (III) (III) (III) (III) (III) (III) (I
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Matches
  Query Match
  Polypeptide having cynomolgus monkey antibody heavy chain variable region or light chain variable region, useful in diagnosing antigen in biological samples, and for treating disease e.g., inflammation and
  07-NOV-2003; 2003US-0517970P
  Example 3; SEQ ID NO 66; 138pp; English
  Human; antibody; interleukin; IL-4; antagonist; receptor; IL-4 receptor; antiarthritic; dermatological; antiulcer; antiinflammatory; cytostatic; antisickling; immunosuppressive; tuberculostatic; ophthalmological;
  Sequence 114
  (AMGE-) AMGEN INC
  26-MAR-2002
   ABB07232 standard; protein; 115
  Anti-IL-4 receptor MAb 12B5 heavy chain variable region.
  Local
  2005-372347/38
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  H
  AEA11010.
  Similarity
   QVQLVQSGGGVVHPGRSLKLSCAGSGFTFS 30
  Shen W,
   Conservative
  (first entry)
   89.6%;
   Jacobsen FW,
   ω
••
   Score 138; DB Pred. No. 1.3e 3; Mismatches
   ₽
   Morris AE,
  ; DB 9;
1.3e-10;
  30
   Allen MJ
  Length 114;
   0
   Gaps
   0
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Homo sapiens

RESULT

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antianemic; antithyroid

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cc antagonist) (1) that binds human II.4 receptor (II.4R), and is capable of conhibiting an II.4-induced biological activity (I) is also useful for cc inhibiting both II.4-induced biological activity and II.-13-induced biological activity and II.-13-induced biological activity and II.-13-induced biological activity in vivo in a human, and for treating septic arthritis conditions such as septic/reactive arthritis, dermatitis herpetiformis, cc unticaria (sepecially chronic idiopathid urticaria), ulcers, gastric cc inflammation, mucosal inflammation, ulcerative colitis, Crohn's disease, characteria especially chronic idiopathid urticaria), ulcers, gastric cc inflammatory bowel disease, other disorders of the digestive system in cc which II.4 plays a role (e.g. II.4-induced inflammation of part of the cg gastrointestinal tract), conditions in which II.4-induced barrier cd disruption plays a role (e.g. conditions) characterized by decreased cc epithelial barrier function in the lung or gastrointestinal tract), cc scleroderma, hypertrophic scarring, whitple's disease, benign prostate chapterialsa, II.4-induced pulmonary conditions, allergic reactions to cmedication, Kawasaki disease, sickle cell disease or crisis, churg-cc strause syndrome, Grave's disease, pre-eclampsia, Sjogren's syndrome, caracter's esophagus, autoimmune uveitis, tuberculosis, nephrosis, cu pemphigus vulgaris or bullous pemphigoid (autoimmune histering contactors also find use as adjuvante to allergy immunotherapy and as caracially when direction the immune rearronae toward a creation adjuvante seasonially when direction in the immune premones toward a
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  Matches
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  26-MAY-2000; 2000US-00579808:
19-SEP-2000; 2000US-00665343:
15-FEB-2001; 2001US-00785934:
01-MAY-2001; 2001US-00847816
   Novel human antibody which binds human interleukin (IL)-4 receptor and is capable of inhibiting IL-4 induced biological activity, functions as IL-4 antagonist and is useful for treating septic arthritis, scleroderma.
  Region
  Region
   Key
  Sequence 115
  vaccine adjuvants, especially when directing the immune response toward a TH1 response would be beneficial in trecting or preventing the disease. The present sequence represents an anti-IL-4 receptor monoclonal antibody
   Claim 4; Page 75-76; 85pp; English.
   WPI; 2002-114332/15.
N-PSDB; ABA94332.
  25-MAY-2001; 2001WO-US017094
  06-DEC-2001.
   Region
   The invention
  Pluenneke
  (IMMV ) IMMUNEX CORP
  present sequence represents an anti-IL-4

o) 12B5 heavy chain variable region
  Similarity
SVQLVQSGGGLVHPGGSLRLSCAGSGFTFS
                                      QVQLVQSGGGVVHPGRSLKLSCAGSGFTFS
  Conservative
  A
   relates to a
  /note=
98. .10
   Location/Qualifiers
  /note=
  . 65
  89.6%;
  "complementarity
  "complementarity determining
  "complementarity
   human antibody (an interleukin (IL)-4
  Score 138; DB 5;
pred. No. 1.4e-10;
3; Mismatches 1
  determining
  determining
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    မ
  Length 115;
  region
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  (CDR)
   (CDR)
  (CDR)
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  Gaps
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ABA11028

ABA11028

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  The invention relates to an isolated polypeptide (I) comprising (a) one CC of 4 sequences: ARA10964, ARA10966, ARA10978 and ARA10978; or ARA10970 or CC ARA10972, and antibody heavy chain variable region. Or (b) ARA10988, and CC antibody light chain variable region. In (I), the antibody heavy chain or Clight chain variable region is a cynomolgus monkey antibody heavy chain or CC region. The antibody heavy chain variable region or cynomolgus monkey antibody light chain variable cregion. Also included are: an isolated polynucleotide (II) CC comprising a sequence encoding (I); an isolated antibody (Al) comprising CC a first polypeptide having SEQ ID No. 30; a method of evaluating (M1) the CC effects of an antibody, comprising introducing into a cynomolgus monkey a comprising the chain variable regions or cynomolgus monkey, and evaluating the effects of the cynomolgus monkey; and evaluating the effects of the chimeric antibody in the cynomolgus monkey, and evaluating the effects of the chimeric antibody in the cynomolgus monkey; an expression vector (V1) comprising (II); a cell CC (III) comprising V1; and a method of producing (I). (Al) further CC comprises a human antibody heavy chain variable region and a mouse antibody light chain variable region. (III) is useful for producing a chimeric antibody, which involves incubating CC antibody light chain variable region and a mouse antibody light chain voriable region. (III) candificant the chimeric antibody. (II) or (Al) further confidencies of the chimeric antibody, which involves incubating CC antison and cancer. The present sequence represents an anti-II-4R confidency chain variable and anti-II-4R confidency chain variable and anti-II-4R confidency chain variable and concer. The present sequence represents an anti-II-4R confidency chain variable and concer. The present sequence represents an anti-II-4R confidency chain variable region.
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  Polypeptide having cynomolgus monkey antibody heavy chain variable region or light chain variable region, useful in diagnosing antigen in biological samples, and for treating disease e.g., inflammation and
  HIV infection; anti-hiv; infection; inflammation; antiinflammatory; cancer; cytostatic; antibody; chimeric antibody; heavy chain variable region; antibody engineering; anti-IL-4R.
   Sequence 115 AA;
  Example 3; SEQ ID NO 70; 138pp; English
  07-NOV-2003; 2003US-0517970P
  04-NOV-2004; 2004WO-US037241.
   26-MAY-2005
   WO2005047325-A2
  Homo sapiens.
   Human anti-IL-4R heavy chain variable region, SEQ ID NO:70
  28-JUL-2005
   AEA11028 standard; protein; 115
  heavy chain variable region used to make the chimeric antibodies
   (AMGE-) AMGEN INC
                                   Local Similarity
  2005-372347/38.
26; Conservative
   Shen W,
  (first entry)
                                   89.6%;
  Jacobsen FW,
Mismatches
                                   Score 138; DB 9;
Pred. No. 1.4e-10;
   Morris AE,
   Allen MJ;
   Indels
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QVQLVQSGGGVVHPGRSLKLSCAGSGFTFS 30

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무
The invention relates to an isolated polypeptide (I) comprising (a) one of 4 sequences: ABA10964, AEA10966, AEA10978, or AEA10970 or CC ABA10972, and antibody heavy chain variable region, or (b)AEA10988, and CC antibody light chain variable region. In (I), the antibody heavy chain or CC variable region or cynomolgus monkey antibody heavy chain or CC region. The antibody heavy chain variable region or light chain variable region or light chain variable region or light chain variable region or light chain variable region or light chain variable region or light chain variable region or light chain variable region or light chain variable region or light chain variable region or light chain variable region or light chain variable region or light chain variable region or light chain variable region or light comprising a sequence encoding (I); an isolated polymucleotide (II) comprising SEQ ID No. 30; a method of evaluating (M1) the effects of an antibody comprising introducing into a cynomolgus monkey and light chain and heavy chain variable regions from a comprising light chain and heavy chain variable regions from a comprise antibody and light chain and heavy chain constant regions from a comprise antibody and light chain and heavy chain constant regions from a comprise and a method of producing (I). (A1) further comprise and a man antibody in the effects of the chimeric antibody in the comprise and a muman antibody heavy chain variable region and a human comprise and a muman antibody light chain variable region and a human comprise antibody. (II) or (A1) further comprise and a muman antibody light chain variable region and a human comprise antibody. (II) or (A1) is useful for diagnosing comprises a human antibody. (III) or (A1) is useful for diagnosing comprises and for treating diseases such as HIV, comprise and the chimeric antibody. (III) or (A1) is useful for diagnosing comprise and comprise and comprise and comprise and comprise and comprise and comprise and comprise and comprise and comprise and comprise and 
  Polypeptide having cynomolgus monkey antibody heavy chain variable region or light chain variable region, useful ih diagnosing antigen in biological samples, and for treating disease e.g., inflammation and
   HIV infection; anti-hiv; infection; inflammation; antiinflammatory; cancer; cytostatic; antibody; chimeric antibody; heavy chain variable region; antibody engineering; anti-IL-4R.
   Aldrich T,
   07-NOV-2003; 2003US-0517970P
   Human anti-IL-4R heavy chain variable region, SEQ ID NO:65
  28-JUL-2005
   AEA11023;
  AEA11023 standard;
  Example 3; SEQ ID NO 65; 138pp; English.
  cancer.
   WPI; 2005-372347/38.
  04-NOV-2004; 2004WO-US037241
   26-MAY-2005
  WO2005047325-A2
  (AMGE-) AMGEN INC.
   Shen W,
  (first entry)
  protein; 115
  Jacobsen FW,
  ⋛
  Morris AE,
  Allen MJ
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   HIV infection; anti-hiv; infection; inflammation; antiinflammatory; cancer; cytostatic; antibody; chimeric antibody; heavy chain variable region; antibody engineering; anti-IL-4R.
  28-JUL-2005
   AEA11027
  Homo sapiens
   Human anti-IL-4R heavy chain variable region, SEQ ID NO:69
  07-NOV-2003; 2003US-0517970P
  04-NOV-2004; 2004WO-US037241
  26-MAY-2005
  (AMGE-) AMGEN INC
  WO2005047325-A2
  26;
  standard; protein; 115 AA
  EVOLVOSGGGLVHPGGSLRLSCAGSGFTFS 30
  Conservative
  (first entry)
  89.6%;
  Jacobsen FW,
   Mismatches
  Score 138; DB 9;
Pred. No. 1.4e-10;
  Allen MJ
   Length 115;
  Indels
  0
  Gaps
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Polypeptide having cynomolgus monkey antibody heavy chain variable region or light chain variable region, useful in diagnosing antigen in biological samples, and for treating disease e.g., inflammation and

WPI; 2005-372347/38. N-PSDB; AEA11013.

Shen W,

Morris AE,

Example 3; SEQ ID NO 69; 138pp; English

The invention relates to an isolated polypeptide (I) comprising (a) one CC of 4 sequences: AEA10964, AEA10966, AEA10978; or AEA10970 or CC AEA10972, and antibody heavy chain variable region. In (I), the antibody heavy chain or CC light chain variable region is a cynomolgus monkey antibody heavy chain or CC variable region or cynomolgus monkey antibody light chain variable region. The antibody heavy chain variable region or light chain variable region is a human or mouse antibody heavy chain variable region or light chain variable region. Also included are: an isolated polymucleotide (II) comprising a sequence encoding (I); an isolated polymucleotide (II) comprising a sequence encoding (I); an isolated antibody (A1) comprising CC after polypeptide having SEQ ID No. 5, 8, 10, 12, 14 or 20, and a comprision of an antibody comprising introducing into a cynomolgus monkey a comprising introducing into a cynomolgus monkey and light chain and heavy chain variable regions from an antibody and light chain and heavy chain constant regions from a cynomolgus monkey, and evaluating the effects of the chimeric antibody in the cynomolgus monkey, and expression vector (V1) comprising (II); a cell (III) comprising V1; and a method of producing (I). (A1) further comprises a human antibody heavy chain variable region and a human comprise antibody light chain variable region and a mouse antibody heavy chain variable region. (III) under conditions suitable for expression of the chimeric antibody, antigen in biological samples, and for treating diseases such as HIV, inflammation and cancer. The present sequence represents an anti-II-4R

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CT The invention relates to an isolated polypeptide (I) comprising (a) one CC 484,0972, and antibody heavy chain variable region, or (b)AEA10988, and CC AEA10972, and antibody heavy chain variable region. In (I), the antibody heavy chain or CC 11ght chain variable region in (I), the antibody heavy chain or CC 11ght chain variable region is a cynomologus monkey antibody heavy chain or CC variable region or cynomologus monkey antibody heavy chain variable region. The antibody heavy chain variable region or 11ght chain variable region is a human or mouse antibody heavy chain variable region or 11ght chain variable region is a human or mouse antibody heavy chain variable region or 11ght CC chain variable region. Also included are: an isolated polymucleotide (II) CC comprising a sequence encoding (I); an isolated antibody (Al) comprising CC a first polypeptide having SEQ ID No. 50; a method of evaluating (M1) the CC effects of an antibody, comprising introducing into a cynomologus monkey, and evaluating the effects of the cynomologus monkey, and evaluating the effects of the cynomologus monkey; an expression vector (V1) comprising (II); a cell CC (III) comprising V1; and method of producing (I). (Al) further CC comprises a human antibody heavy chain variable region and a human antibody light chain variable region antibody heavy chain variable region and a human antibody light chain variable region antibody heavy chain variable region and a human antibody light chain variable region and a human antibody light chain variable region and a human antibody light chain variable region and a human antibody light chain variable region and a human antibody light chain variable region and a human antibody light chain variable region and a human antibody light chain variable region and a human antibody light chain variable region and antibody light chain variable region and antibody light chain variable region and antibody light chain variable region and antibody light chain variable region and antibody light chain variable reg
  Matches
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  heavy chain variable region used to make the chimeric antibodies of the
   cancer; cytostatic; antibody; chimeric antibody; heavy chain variable region; antibody engineering; anti-IL-4R
   28-JUL-2005
  Sequence 115
   Example 3; SEQ ID NO 67; 138pp; English
  Polypeptide having cynomolgus monkey antibody heavy chain variable region or light chain variable region, useful in diagnosing antigen in biological samples, and for treating disease e.g., inflammation and
  WPI; 2005-372347/38.
   07-NOV-2003; 2003US-0517970P
  04-NOV-2004; 2004WO-US037241
   26-MAY-2005
   Homo
   HIV infection; anti-hiv; infection; inflammation; antiinflammatory;
   Human anti-IL-4R heavy chain variable region, SEQ ID NO:67
  AEA11025;
   ABA11025
  invention.
   cancer.
   Aldrich T,
   (AMGE-) AMGEN INC
   WO2005047325-A2
  26;
   Similarity
   AEA11011.
  standard; protein; 115 AA.
  EVOLVOSGGGLVHPGGSLRLSCAGSGFTFS 30
   Shen W,
  Conservative
  ₹
  (first entry)
  89.6%;
   Jacobsen FW,
  ω
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  Score 138; DB Pred. No. 1.4e 3; Mismatches
   Morris AE,
   138; DB 9;
No. 1.4e-10;
  DΒ
  9;
   Allen MJ
  Length 115
  Indels
  0
  Gaps
   0
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The invention relates to an isolated polypeptide (I) comprising (a) one of 4 sequences: AEA10964, AEA10966, AEA10968 and AEA10978, or AEA10970 or AEA10972, and antibody heavy chain variable region, or (b) AEA10988, and antibody light chain variable region. In (I), the antibody heavy chain or light chain variable region is a cynomolgus monkey antibody heavy chain or variable region. The antibody heavy chain variable region or light chain variable region or light chain variable region is a human or mouse antibody heavy chain variable region or light chain variable region. Also included are: an isolated polymucleotide (II) comprising a sequence encoding (I); an isolated polymucleotide (II) a first polypeptide having SEQ ID No. 6, 8, 10, 12, 14 or 20, and a second polypeptide having SEQ ID No. 30; a method of evaluating (M1) the second polymental second polymental second polymental second polymental second second polymental second second polymental second second polymental second second second polymental second s
   Query Match
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Matches 26
  is useful for producing a chimeric antibody, which involves incubating (III) under conditions suitable for expression of the chimeric antibody, and isolating the chimeric antibody. (I) or (A) is useful for diagnosing antigen in biological samples, and for treating diseases such as HIV, inflammation and cancer. The present sequence represents an anti-IL-4R heavy chain variable region used to make the chimeric antibodies of the
   Polypeptide having cynomolgus monkey antibody heavy chain variable region or light chain variable region, useful in diagnosing antigen in biological samples, and for treating disease e.g., inflammation and
   Homo sapiens
   HIV infection; anti-hiv; infection; inflammation; antiinflammatory; cancer; cytostatic; antibody; chimeric antibody;
  Human anti-IL-4R heavy chain variable region, SEQ ID NO:73
  Sequence 115 AA;
   Example 3; SEQ ID NO 73; 138pp; English
   (AMGE-) AMGEN INC
  07-NOV-2003; 2003US-0517970P
   04-NOV-2004; 2004WO-US037241
   WO2005047325-A2
   heavy chain variable region; antibody engineering; anti-IL-4R
  28-JUL-2005
   AEA11031 standard; protein; 115
   2005-372347/38
   26;
  1 QVQLVQSGGGVVHPGRSLKLSCAGSGFTFS
:|||||||||||||||
   AEA11017.
   Similarity
  EVQLVQSGGGLVHPGGSLRLSCAGSGFTFS 30
  Shen W,
  Conservative
  (first entry)
  89.6%;
  Jacobsen FW,
  ; Score 138; DB; Pred. No. 1.4e: 3; Mismatches
  Morris AE,
  30
   .4e-10;
  멂
  9
   Allen MJ;
  Length 115;
   Indels
  Gaps
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   맑
  the cynomolgus monkey; an expression vector (V1) comprising (II); a cell (III) comprising V1; and a method of producing (I). (A1) further comprises a human antibody heavy chain variable region and a human antibody light chain variable region or a mouse antibody heavy chain variable region and a mouse antibody light chain variable region. (III) is useful for producing a chimeric antibody, which involves incubating (III) under conditions suitable for expression of the chimeric antibody, and isolating the chimeric antibody. (I) or (A1) is useful for diagnosing antigen in biological samples, and for treating diseases such as HIV, inflammation and cancer. The present sequence represents an anti-IL-R heavy chain variable region used to make the chimeric antibodies of the
The invention relates to an isolated polypeptide (I) comprising (a) one of 4 sequences: ABA10964, ABA10966, ABA10978 and ABA10970 or ABA10970 or ABA10972, and antibody heavy chain variable region, or (b) ABA10988, and antibody light chain variable region. In (I), the antibody heavy chain or light chain variable region is a cynomolgus monkey antibody heavy chain or variable region or cynomolgus monkey antibody light chain variable region. The antibody heavy chain variable region is a human or mouse antibody heavy chain variable region is a human or mouse antibody heavy chain variable region or light chain variable region. Also included are: an isolated polynucleotide (II) comprising a sequence encoding (II) an isolated antibody (AI) comprising a first polypeptide having SEQ ID No. 6, 8, 10, 12, 14 or 20, and a
  Polypeptide having cynomolgus monkey antibody heavy chain variable region or light chain variable region, useful in diagnosing antigen in biological samples, and for treating disease e.g., inflammation and
  HIV infection; anti-hiv; infection; inflammation; antiinflammatory; cancer; cytostatic; antibody; chimeric antibody; heavy chain variable region; antibody engineering; anti-IL-4R.
  Sequence 115
   Example 3;
  N-PSDB;
   Aldrich T,
   07-NOV-2003; 2003US-0517970P
   04-NOV-2004; 2004WO-US037241.
  26-MAY-2005
  WO2005047325-A2
   Human anti-IL-4R heavy chain variable region,
   28-JUL-2005
  AEA11022 standard; protein; 115
  Homo sapiens.
  AEA11022;
   (AMGE-) AMGEN INC
   Local Similarity
  2005-372347/38.
)B; AEA11008.
   سا
   EVQLVQSGGGLVHPGGSLRLSCAGSGFTFS
   QVQLVQSGGGVVHPGRSLKLSCAGSGFTFS 30
   SEQ ID NO 64; 138pp; English
   Shen W,
   Conservative
  B
   (first entry)
   89.6%;
   Jacobsen
   ω
•
   Score 138; DB 9;
Pred. No. 1.4e-10;
   Œ,
   Mismatches
  8
   Morrib AE,
   30
   SEQ ID NO:64
   Allen MJ
   Length 115;
   0,
   Gaps
   0
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RESULT 15
AEA11020
ID AEA111
XX AEA11
XX AEA11
XX AEA11
XX AEA11
XX HUWAN
XX HUWAN
XX HOMO
DE HUMAN
XX HOMO
DE HOMO
DE HOMO
XX HOMO
DY WO200
XX HOMO
XX HOMO
YX WO21
PR 04-NU
XX O7-NU
PR 07-NU
PR N-PSI
XX N-PSI
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  2020202020202020202020
   밁
   ई
  CC second polypeptide having SEQ ID No. 30; a method of evaluating (M1) the effects of an antibody, comprising introducing into a cynomolgus monkey a CC chimeric antibody comprising light chain and heavy chain variable regions CC from an antibody and light chain and heavy chain constant regions from a CC cynomolgus monkey, and evaluating the effects of the chimeric antibody in the cynomolgus monkey, an expression vector (V1) comprising (II); a cell CC (III) comprising V1; and a method of producing (I). (A1) further CC comprises a human antibody heavy chain variable region and a human CC antibody light chain variable region or a mouse antibody heavy chain (III) is useful for producing a chimeric antibody, which involves incubating CC is useful for producing a chimeric antibody, which involves incubating CC interest of the chimeric antibody. (I) or (A1) is useful for diagnosing CC antigen in biological samples, and for treating diseases such as HIV, inflammation and cancer. The present sequence represents an anti-IL-4R chimeric chain variable region used to make the chimeric antibodies of the
   Best Local
  Query Match
  Matches
  HIV infection; anti-hiv; infection; inflammation; antiinflammatory; cancer; cytostatic; antibody; chimeric antibody; heavy chain variable region; antibody engineering; anti-IL-4R.
   Sequence 115 AA;
   07-NOV-2003; 2003US-0517970P
  04-NOV-2004; 2004WO-US037241
   26-MAY-2005
  WO2005047325-A2
  Human anti-IL-4R heavy chain variable region, SEQ ID NO:62.
  28-JUL-2005
   AEA11020,
  AEA11020 standard; protein; 115 AA
   Polypeptide having cynomolgus monkey antibody heavy chain variable region or light chain variable region, useful in diagnosing antigen in biological samples, and for treating disease e.g., inflammation and
  N-PSDB; AEA11006.
   WPI; 2005-372347/38.
  Aldrich T,
  AMGE-) AMGEN INC
   ۳
  Similarity
   QVQLVQSGGGVVHPGRSLKLSCAGSGFTFS 30
   EVQLVQSGGGLVHPGGSLRLSCAGSGFTFS
  Shen W,
   Conservative
  (first entry)
  89.6%;
   Jacobsen FW,
   ω
--
  Score 138; DB 9;
Pred. No. 1.4e-10;
   Mismatches
   Morris AE, Allen MJ;
  30
  9; Length 115;
   Indels
   0,
   Gaps
```

The invention relates to an isolated polypeptide (I) comprising (a) one of 4 sequences: AEA10964, AEA10966, AEA10968 and AEA10978; or AEA10970 or AEA10972, and antibody heavy chain variable region, or (b) AEA10988, and antibody heaving chain or light chain variable region. In (I), the antibody heavy chain or light chain variable region is a cynomolgus monkey antibody heavy chain variable region or cynomolgus monkey antibody light chain variable

Example 3; SEQ ID NO 62; 138pp; English.

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CC region. The antibody heavy chain variable region or light chain variable credion is a human or mouse antibody heavy chain variable region or light CC chain variable region. Also included are: an isolated polynucleotide (II) CC comprising a sequence encoding (I); an isolated antibody (AI) comprising CC a first polypeptide having SEQ ID No. 6, 8, 10, 12, 14 or 20, and a CC second polypeptide having SEQ ID No. 30; a method of evaluating (MI) the CC effects of an antibody, comprising introducing into a cynomolgus monkey a CC cynomolgus monkey and light chain and heavy chain variable regions from a notibody and light chain and heavy chain constant regions from a CC cynomolgus monkey; an expression vector (V1) comprising (II); a cell CC (III) comprising V1; and a method of producing (I). (A1) further CC comprises a human antibody heavy chain variable region and a human cC comprises a human antibody heavy chain variable region and a human cC comprises a human antibody heavy chain variable region and a human cC comprises a human antibody heavy chain variable region and a human cC comprises a human antibody heavy chain variable region and a human cC comprises a human antibody heavy chain variable region and a human cC comprises a human antibody heavy chain variable region and a human cC comprises a human antibody. (II) or (A1) is useful for diagnosing confidence conditions suitable for expression of the chimeric antibody, cand isolating the chimeric antibody. (I) or (A1) is useful for diagnosing confidence conditions suitable region and a method of producing dependence represents an anti-II-4R heavy chain variable region used to make the chimeric antibodies of the involve chain variable region used to make the chimeric antibodies of the involve chain variable region used to make the chimeric antibodies of the involve chain variable region used to make the chimeric antibodies of the involve chain variable region used to make the chimeric antibodies of the involve chain variable region used to make the chimeric antibod
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Sequence 115 A,

0

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밁
  Matches
  Query Match
   Local
                          1 QVQLVQSGGGVVHPGRSLKLSCAGSGFTFS 30
   Similarity
EVQLVQSGGGLVHPGGSLRLSCAGSGFTFS
  Conservative
  89.6%; Score 138; DB 9;
86.7%; Pred. No. 1.4e-10;
tive 3; Mismatches 1
  Length 115;
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  Gaps
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Search completed: November 21, Job time: 41.1648 secs 2005, 12:20:03 This Page Blank (uspto)

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Minimum DB
Maximum DB
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Maximum Match 100%
Listing first 45 summaries
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  1 QVQLVQSGGGVVHPGRSLKLSCAGSGFTFS 30
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3: pir3:*
4: pir4:*
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   GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
  2005, 11:49:36; Search time 7.30769 Seconds (without alignments)
394.995 Million cell updates/sec
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

| 29                 | 28             | 27          | 26          | 25            | 24          | 23            | 22             | 21     | 20          | 19             | 18          | 17          | 16            | 15     | 14            | 13     | 12            | 11            | 10            | 9      | 8             | 7              | o             | տ           | 4              | ω             | N              | 1      | Result<br>No.  |
|--------------------|----------------|-------------|-------------|---------------|-------------|---------------|----------------|--------|-------------|----------------|-------------|-------------|---------------|--------|---------------|--------|---------------|---------------|---------------|--------|---------------|----------------|---------------|-------------|----------------|---------------|----------------|--------|----------------|
| 130                | 131            | 131         | 131         | 134           | 134         | 134           | 134            | 134    | 134         | 134            | 134         | 134         | 134           | 134    | 134           | 134    | 134           | 134           | 134           | 134    | 134           | 134            | 134           | 134         | 134            | 137           | 137            | 138    | Score          |
| 84.4               | 85.1           | 85.1        | 85.1        | 87.0          |             | 87.0          | ?              |        | .7          | 87.0           | .7          | 7.          | 7.            | •      | •             | .7     | 7.            | 7.            | .7            | 87.0   | .7            | ?              | 7             | .7          | .7             | 89.0          | .9             | ۰.     | Query<br>Match |
| 97                 | 122            | 114         | 97          | 139           | 137         | 134           | 133            | 133    | 132         | 130            | 130         | 128         | 122           | 122    | 122           | 121    | 120           | 119           | 119           | 118    | 113           | 113            | 98            | 98          | 94             | 121           | 119            | 116    | Length         |
| N                  | -              | N           | N           | N             | N           | N             | N              | N      | Ŋ           | N              | N           | N           | N             | N      | N             | N      | N             | N             | N             | N      | N             | N              | N             | N           | N              | N             | μ              | N      | B .            |
| S26886             | мзниам         | S46390      | S44115      | S31674        | 831701      | S31679        | A49028         | S31590 | 831603      | B60014         | S31601      | S48797      | 831119        | E36005 | 831117        | G36005 | 831112        | F36005        | 831111        | S31116 | S38490        | S57410         | S29546        | PL0116      | PL0120         | S19666        | GIHUNI         | 812557 | ID             |
| Ig heavy chain V r | heavy chain V- | heavy chain | heavy chain | heavy chain V | heavy chain | heavy chain V | heavy chain V- | heavy  | heavy chain | heavy chain pr | heavy chain | heavy chain | heavy chain - | heavy  | heavy chain - |        | heavy chain - | heavy chain V | heavy chain - | heavy  | heavy chain - | heavy chain V- | heavy chain V | heavy chain | heavy chain V- | heavy chain V | heavy chain V- | heavy  | Description    |

| 45     | 44     | 43     | 42     | 41     | 40     | 39     | 38     | 37     | 36     | S      | 34     | 33     | 32     | 31     | 30     |
|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|
| 125    | 125    | 125    | 125    | 125    | 126    | 126    | 127    | 127    | 127    | 127    | 128    | 128    | 129    | 130    | 130    |
| 81.2   | 81.2   | 81.2   | 81.2   | 81.2   | 81.8   | 81.8   | 82.5   | 82.5   | 82.5   | 82.5   | 83.1   | 83.1   | 83.8   | 84.4   | 84.4   |
| 127    | 126    | 122    | 121    | 97     | 134    | 123    | 140    | 135    | 134    | 98     | 151    | 115    | 96     | 127    | 120    |
| N      | ۳      | ۲      | ۲      | N      | N      | N      | ผ      | N      | N      | N      | N      | N      | N      | N      | ผ      |
| S19878 | GIHUKL | M3HUGA | G1HUHL | S26885 | S31699 | S38493 | S70442 | 831598 | S31688 | S29543 | A60943 | S57445 | PH0873 | S38489 | 836273 |
|        |        |        |        |        |        |        | •      |        |        |        |        |        |        |        |        |
| ΕĪ     | ρī     | ĒĪ     | ΕĪ     | ξĪ     | ξĪ     | EI     | ξĨg    | EI     | ξīg    | PI     | ΕĪ     | 51     | Бт     | 19     | Įg     |
| neavy  | neavy  | neavy  | neavy  | neavy  | neavy  | neavy  | neavy  | neavy  | neavy  | neavy  | neavy  | neavy  | neavy  | neavy  | heavy  |
| cnain  | cnain  | chain  | cnain  | chain  | chain  | chain  | chain  | chain  | cnain  | cnain  | chain  | cnain  | chain  | chain  | chain  |
| 4      |        | : <    | : <    | : <    | ; <    | : :    | pre    | \<br>H | : <    | : <    | pre    | \ -C   | : <    | : :    | ۲ H    |
|        |        |        |        |        |        |        |        |        |        |        |        |        |        |        |        |

### ALIGNMENTS

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A,Cross-references: UNIPROT:P01770; UNIPARC:UPI000012CEFA
A,Note: the sequence of the gamma-1 C region of this chain
R, preker, L.; Schwarz, J.; Reichel, W.; Hilsdhmann, N.
Hoppe-Seyler's Z. Physiol. Chem. 357, 1515-1940, 1976
A, Title: Rule of antibody structure. The primary structure enbromide cleavage products, and the disulfide bridges.
A,Reference number: A91667, MUID:77070267; PMID:1002129
A,Contents: annotation; disulfide bond
C;Comment: This chain was isolated from an IgG1 myeloma pro
  Ig heavy chain V-III region (Nie) - human
C;Species: Homo sapiens (man)
C;Species: 23-Oct-1981 #sequence_revision 23-Oct-1981 #text_change 09-Jul-2004
C;Accession: A91668; A02053
C;Accession: A91668; A02053
R;Ponstingl, H.; Hilschmann, N.
Hoppe-Seyler's Z. Physiol. Chem. 357, 1571-1604, 1976
A;Title: Die Primaerstruktur eines monoklonalen IgG1-Immunglobulins (Myelomprotein Nie),
  Ig heavy chain - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 21-Nov-1993 #sequence_revision 13-Mar-1997 #text_change 21-Jan-2000
C;Date: 21-Nov-1993
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   Ś
   A;Cross-references: UNIPARC:UPI000013DE0E
C;Superfamily: immunoglobulin V region; immunoglobulin homology
F;34-116/Domain: immunoglobulin homology <IMM>
  A;Reference number: S12557; MUID:90316107; PMID:2114977 A;Accession: S12557
   A; Title: Organization of variable region segments of the human immunoglobulin heavy chair
   C;Accession: S12557
R;Matsuda, F.; Shin, E.K.; Hirabayashi, Y.; Nagaoka, H.; Yoshida, M.C.; Zong, S.Q.; Honjc EMBO J. 9, 2501-2506, 1990
  A; Molecule type: DNA
A; Residues: 1-116 < MAT>
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Pred. No. 1.9e-11;
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   49
  ۲.
  Indels
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   of monoclonal IgG1 immunogloby
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  0
  Gaps
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C!Species: Homo saplens (man)
C!Species: Homo saplens (man)
C!Date: 22-Jan-1993 #sequence_revision 22-Jan-1993 #text_change 20-Jun-2000
C!Accession: 819666
R!Marks, J.D.; Hoogenboom, H.R.; Bonnert, T.P.; McCafferty, J.; Griffiths, A.D.; Winter, J. Mol. Biol. 222, 581-597, 1991
A;Title: By-passing immunization. Human antibodies from V-gene libraries displayed on ph A;Reference number: S19663; MUID:92085276; PMID:1748994
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A;Grose references: GDB:128528; OMIM:147070
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C;Keywords: heterotetramer; immunoglobulin; pyroglutamic acid
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F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status ex
F;22-96/Disulfide bonds: #status experimental
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C;Date: 07-Un-1990 #sequence_revision 07-Jun-1990 #text_change 31-Dec-
C;Accession: PLO120
R;Bird, J.; Galill, N.; Link, M.; Stites, D.; Sklar, J.
J. Exp. Med. 168, 229-245, 1988
A;Title: Continuing rearrangement but absence of somatic hypermutation
A;Reference number: PLO116; MUID:88286083; PMID:2840480
A;Accession: PLO120
   A;Cross-references: UNIPROT:Q8WUK1; UNIPROT:Q9UL93; UNIPARC:UPI0000176A2C A;Experimental source: B cells from patient TD with acute lymphoblastic leather the sequence shows the V region (TD-Vo) from a nonproductive DNA;C;Superfamily: immunoglobulin homology
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Best Local :
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90.0%; Pred. No. 2.6e-11;
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   Score 134; DB 2;
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  Length 121
  Length 94;
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  0
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  Gaps
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   in immunoglobulin
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   PIC
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A,Experimental Source: B cells from patient AW with acute lymphoblastic leukemia, ALL A,Note: the sequence shows the V region (AW Vx) from a nonproductive DNA rearrangement for prolinson, i.M.; Walter, G.; Marks, J.D.; Llewelyn, M.B.; Winter, G. J. Mol. Biol. 227, 776-798, 1922

A,Title: The repertoire of human germline V(H) sequences reveals about fifty groups of V. A,Reference number: $26885; MUID:93021117; PMID:1404388

A;Accession: $26892

A;Status: preliminary A,Accession: $26895

A;Coss.-references: UNIPARC:UPI000031F3A; EMBL:Z12349; NID:g32918; PIDN:CAA78219.1; PID: C;Superfamily: immunoglobulin V region; immunoglobulin homology C;Keywords: acute lymphoblastic leukemia; heterotetramer; immunoglobulin F;15-98/Domain: immunoglobulin v region; immunoglobulin homology C;Keywords: acute lymphoblastic leukemia; heterotetramer; immunoglobulin F;15-98/Domain: immunoglobulin v region; immunoglobulin homology C;Keywords: acute lymphoblastic leukemia; heterotetramer; immunoglobulin F;15-98/Domain: immunoglobulin homology 
  A;Cross-references: UNIDARC:UDI000002DD16; EMBL:Z17394; NID:g32843; PIDN:CAA78997.1; PID: A;Note: designated COS-8
R;Tomlinson, I.M.; Walter, G.; Marks, J.D.; Llewelyn, M.B.; Winter, G.
J. Mol. Biol. 227, 776-798, 1992
A;Title: The repertoire of human germline V(H) sequences reveals about fifty groups of V-A;Reference number: S26885; MUID:93021117; PMID:1404388
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C;Species: Homo sapiens (man)
C;Date: 07-Jan-1994 #sequence_revision 17-Nov-1995 #text_c
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C;Species: Homo sapiens (man)
C;Date: 07-Jun-1990 #sequence_revision 07-Jun-1990 #text_change
C;Accession: Pi0116; S26892
   A; Molecule type: DNA
A; Residues: 1-98 < TOM>
   R;Bird, J.; Galili, N.; Link, M.; Stites, D.; Sklar, J. Exp. Med. 168, 229-245, 1988
  A; Molecule type: DNA
A; Residues: 1-98 < TO2>
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  A, Cross-references: UNIPARC: UPI0000031F3A
   Best Loc
Matches
Matches
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  Local
   Local
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  Similarity
  Similarity
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Conservative
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86.7%;
  87.0%;
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Score 134; DB
Pred. No. 5.5e
2; Mismatches
2
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Pred. No. 5.5e-11;
  Mismatches
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   DB 2;
.5e-11;
   #text_change
  Length 98
  Length 98
  23-Jul-1999
   23-Jul-1999
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   Gaps
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C;Accession: S57410
R;Paterson, G.; Wilson, G.; Kennedy, P.G.E.; Willison, H.J.
submitted to the EMBL Data Library, June 1995
A;Description: Analysis of anti-GM1 ganglioside IgM antibod
A;Reference number: S57408
A;Accession: S57408
A;Accession: S57408
   R;Marks, J.D.; Ouwehand, W.H.; Bye, J.M.; Finnern, R.; Gorick, B.D.; Voak, D.; submitted to the EMBL Data Library, June 1993
A;Description: Human antibody fragments specific for human blood group antigens
   A;Introns: 99/2
C;Superfamily: immunoglobulin V region; immunoglobulin homology
F;15-98/Domain: immunoglobulin homology <IMM>
   A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-113 <MAR>
   문
  ई
C;Species: Homo sapiens (man)
C;Date: 02-Dec-1993 #sequence_revision 26-May-1995 #text_change 31-Dec-2004
C;Date: 02-Dec-1993 #sequence_revision 26-May-1995 #text_change 31-Dec-2004
C;Accession: S31116
R;Raaphorst, F.M.; Timmers, E.; Kenter, M.J.H.; van Tol, M.J.D.; Vossen, J.I.
Bur. J. Immunol. 22, 247-251, 1992
A;Title: Restricted utilization of germ-line V(H)3 genes and short diverse A;Reference number: S31104; MUID:92111633; PMID:1730252
  S
   8
  A, Reference number: S38488
  Ig heavy chain -
  뭐
   밁
  Ig heavy chain - human
   Cross-references: UNIPARC:UPI0000116549; EMBL:Z23030; NID:g414027; Superfamily: immunoglobulin V region; immunoglobulin homology Reywords: heterotetramer; immunoglobulin
  g heavy chain - human (fragment)
;Species: Homo sapiens (man)
;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 23-Jul-1999
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   Status: preliminary
  Date: 10-Oct-1995 #sequence_revision 17-Nov-1995 #text_change
  ;Species: Homo sapiens (man)
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  Query Match
   heavy chain V-J region - human (fragment)
   Local Similarity
   26;
  26;
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   QVQLVESGGGVVQPGRSLRLSCAASGFTFS
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  Conservative
  87.0%;
   87.0%;
86.7%;
   B.; Kenter, M.J.H.; van Tol, M.J.D.; Vossen, J.M.; Schuurmar
   2
  Score 134; DB 2;
Pred. No. 6.4e-11;
   Score 134; DB 2;
Pred. No. 6.4e-11;
   Mismatches
  Mismatches
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  Length 113;
   Length 113
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  23-Jul-1999
  group antigens from a
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  Gaps
  PIDN: CAA80565.1;
   PIDN: CAA61144.1;
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A;Note: the nucleotide sequence was submitted C;Superfamily: immunoglobulin homology C;Superfamily: immunoglobulin homology <IMM> P;15-98/Domain: immunoglobulin homology <IMM>
  A;Gene: GDB:IGH@; IGHDY1
A;Cross-references: GDB:118731; OMIM:146910
A;Map position: 14q32.33-14q32.33
C;Superfamily: immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;15-98/Domain: immunoglobulin homology <IM
   C;Accession: F36005
R;Schroeder Jr., H.W.; Wang, J.Y.
Proc. Natl. Acad. Sci. U.S.A. 87, 6146-6150, 1990
Proc. Natl. Acad. Sci. U.S.A. 87, 6146-6150, 1990
A;Title: Preferential utilization of conserved immunoglobulin heavy chain variable gene A;Reference number: A36005; MUID:90349571; PMID:2117273
  RESULT 11
F36005
  C;Superfamily: :
C;Keywords: hete
F;15-98/Domain:
  A;Cross-references: UNIPARC:UP10000176DC2; EMBL:X62959
A;Note: the nucleotide sequence was submitted to the EMBL Data 1
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
  Bur. J. Immunol. 22, 247-251, 1992
A;Title: Restricted utilization of germ-line V(H)3 genes and A;Reference number: S31104; MUID:92111633; PMID:1730252
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R;Raaphorst, F.M.; Timmers, B.; Kenter, M.J.H.; van Tol, M.J.D.; Vossen, J.M.; Schuurman,
  C;Date: 02-Dec-1993 #sequence_revision 26-May-1995 #text_change 17-Mar-1999 C;Accession: S31111
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A;Note: the nucleotide sequence was submitted to the EMBL Data Library, October 1991
   A; Molecule type: mRNA
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C;Species: Homo sapiens (man)
C;Date: 21-Dec-1990 #sequence_revision 21-Dec-1990 #text_change 31-Dec-2004
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   A; Accession: S31111
  A;Residues: 1-119 <RAA>
   Ig heavy chain -
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  A; Accession: F36005
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  Matches
  Query Match
   Query Match
Best Local
                        Query Match
   Local
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  Similarity
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2; Mismatches 2
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2; Mismatches 2
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Pred. No. 6.7e-11;
   ^IMY>
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   30
  2
  Length 119;
  EMBL Data Library, October 1991
   Length
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  Indels
  Indels
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Best Local Similarity

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Ig heavy chain V region (M74) - human
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C;Date: 21-Dec-1990 #sequence_revision 21-Dec-1990 #text_change 31-Dec-2004
C;Accession: G36005
R;Schroeder Jr., H.W.; Wang, J.Y.
Proc. Natl. Acad. Sci. U.S.A. 87, 6146-6150, 1990
A;Title: Preferential utilization of conserved immunoglobulin heavy chain variable gene A;Reference number: A36005; MUID:90349571; PMID:2117273
A;Accession: G36005
   A;Gene: GDB:IGH@; IGHDY1
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G;Accession: S31112

Finmers, E.; Kenter, M.J.H.; van Tol, M.J.D.; Vossen, J.M.; Schuurman R;Raaphorst, F.M.; 22, 247-251, 1992

Bur. J. Immunol. 22, 247-251, 1992

A;Title: Restricted utilization of germ-line V(H)3 genes and short diverse third complem A;Reference number: S31104; MUID:92111633; PMID:1730252

A;Accession: S31112

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A;Note: the nucleotide sequence was submitted to the EMBL Data Library,
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
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C;Species: Homo sapiens (man)
C;Date: 02-Dec-1993 #sequence_revision 26-May-1995 #text_change 17-Mar-1999
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A;Molecule type: mRNA
A;Residues: 1-121 <SCH>
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  र्
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  A;Molecule type: mRNA
A;Residues: 1-120 <RAA>
   Query Match
Best Local Similarity
Matches 26; Conserv
  Query Match 87.0%;
Best Local Similarity 86.7%;
Matches 26; Conservative
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Date: 02-Dec-1993 #sequence_revision 26-May-1995 #text_change 17-Mar-1999
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  30
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A;Gene: GDB:IGH@; IGHDY1
A;Cross-references: GDB:118731; OMIM:146910
A;Map position: 14q32.33-14q32.33
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
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(;Accession: B36005
R;Schroeder Jr., H.W.; Wang, J.Y.
Proc. Natl. Acad. Sci. U.S.A. 87, 6146-6150, 1990
A;Title: Preferential utilization of conserved immunoglobulin heavy chain variable
A;Reference number: A36005; MUID:90349571; PMID:2117273
   C;Accession: S31117

R;Rasphorst, F.M.; Timmers, E.; Kenter, M.J.H.; van Tol, M.J.D.; Vossen, J.M.; Schuur R;Rasphorst, F.M.; 247-251, 1992

A;Title: Restricted utilization of germ-line V(H)3 genes and short diverse third compares and short diverse third compares in the second second sequence for shown; s1117

A;Status: preliminary; nucleic acid sequence hot shown; translation not shown A;Status: preliminary; nucleic acid sequence hot shown; translation not shown A;Molecule type: mRNA
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A;Note: the nucleotide sequence was submitted to the EMBL Data Library, October 1991
C;Superfamily: immunoglobulin v region; immunbglobulin homology
C;Keywords: heterotetramer; immunoglobulin
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   RESULT
E36005
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re greater than or equal to the score of the result being printed,
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   length:
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2: uniprot_trembl:*
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1 homo sapien
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2 homo sapien
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0 homo sapien
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SNR; P01770; 1-119.

GO; GO:0005576; C:extracellular region; NAS.

GO; GO:0005823; F:antigen binding; NAS.

InterPro; IPR0037110; Ig-like.

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SMART; SM00406; IGV; 1.

PROSITE; PS50835; IG LIKE; 1.

PROSITE; PS50835; IG LIKE; 1.

Direct protein sequencing; Immunoglobulin domain; Immunoglobulin V region; Pyrrolidone carboxylic acid DOMAIN

1 112 Ig-like.
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P01770;
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  STANDARD;
   PRT;
  ALIGNMENTS
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|             | 120<br>119<br>119<br>119<br>118<br>118<br>118<br>118<br>118<br>118<br>118                                                                                                                                                                                                  |
|-------------|----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
|             | 77.3<br>77.3<br>77.3<br>77.3<br>76.6<br>76.6<br>76.6<br>76.6                                                                                                                                                                                                               |
|             | 479<br>114<br>519<br>112<br>115<br>115<br>116<br>117<br>117<br>119<br>499                                                                                                                                                                                                  |
|             | 222                                                                                                                                                                                                                                                                        |
| AT.TGNMENTS | QSPQK9 RAT HV36 MÖUSB HV3B-HUMAN QSHCC1 HUMAN HV3D HUMAN HV3B HUMAN HV3C HUMAN HV3C HUMAN HV3C HUMAN HV3C HUMAN HV3C HUMAN QSPSZLB MOUSB HV3M HWOUSB GSPSZLB MOUSB QSPSZLB MOUSB QSPSZLB MOUSB                                                                             |
|             | Q5pqk9 rattus norv p18527 mus musculu p18527 mus musculu p101763 homo sapien Q6n092 homo sapien Q9hcc1 homo sapien p01765 homo sapien p01779 homo sapien p01774 homo sapien p18526 mus musculu p01774 homo sapien Q65z18 mus musculu Q8n5k4 homo sapien Q6pj95 homo sapien |

#### MEDLINE-77070367; PubMed=1002129; Dreker L., Schwarz J., Reichel W., Hilschmann N.; Rule of antibody structure. The primary structure of a monoclonal igg1 immunoglobulin (myeloma protein Nié), I: purification and characterization of the protein, the L- and H-chains, the cyanogen bromide cleavage products, and the disulfide bridges."; Hoppe-Seyler's Z. Physiol. Chem. 357:1515-1540(1976). 21-JUL-1986 (Rel. 01, Created) 21-JUL-1986 (Rel. 01, Last sequence up 10-MAY-2005 (Rel. 47, Last annotation Ig heavy chain V-III region NIE. Homo sepiens (Human). This Swiss-Prot entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation - the European Bioinformatics Institute. There are no restrictions on its MEDLINE-77070269; PubMed-826475; Ponsting1 H., Hilschmann N.; Ponsting1 H., Hilschmann N.; Ponsting1 H., Hilschmann N.; Ponsting1 H. Hilschmann N.; The primary structure of a monoclonal right rule of antibody structure. The primary structure of the chymotryptic peptides of the H-chain, alignment of the tryptic peptides and discussion of the complete structure."; Hoppe-Seyler's Z. Physiol. Chem. 357:1571-1604(1976). Eukaryola; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia, Eutheria; Euarchontoglires; Primates; Catarrhini, Hominidae; protein. SIMILARITY: Contains 1 Ig-like (immunoglobulin-like) domain. in no way modified and this statement ) updáte) on update) A 18

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  RC TISUE-Primary B-Cells;

RC TISUE-Primary B-Cells;

RS HEDLINE-2388257, pubMed-12477932; DOI=10.1073/pnas.242603899;

RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

RA Klausner R.D., Collins F.S., Wagner L., Shammen C.M., Schuler G.D.,

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RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

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RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

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RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

VII Bosak S.A., McEwan P.J., McKernan M., Madan A., Rodrigues S., Sanchez A.,

RA Niltalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

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Ra Holtseye R.M., Touchman J.W., Green E.D., Dickson M.C.,

Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,

"Generation and initial analysis of more than 15,000 full-length human

and mouse cDNA sequences.",

Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903 (2002).
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PubMed=2117273;

Schroeder H.W. Jr, Wang J.Y.;

Schroeder H.W. Jr, Wang J.Y.;

*Preferential utilization of conserved immunoglobulin heavy

*Preferential utilization of the second immunoglobulin heavy

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variable gene segments during human fetal life.";

variable gene segments during human fetal life.";
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PubMed=1383695; DOI=0.1016/0161-5890(92)90173-U;

PubMed=1383695; DOI=0.1016/0161-5890(92)90173-U;

Cuisinier A.M., Funcoux F., Fougereau M., Tonnelle C.;

TIGM kappa/lambda BBV human B cell clone: an early step of

"IGM kappa/lambda BBV human B cells or a distinct B lineage?";
            NUCLEOTIDE SEQUENCE
  differentiation of fetal B cells or Mol. Immunol. 29:1363-1373(1992).
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96
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Last sequence update)
Last annotation update)
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   PubMed=253855; DOI=10.1084/jem.169.4.1391;
Nickerson K.G., Berman J., Glickman E., Chess L., Alt F.W.;
"Early human IgH gene assembly in Epstein-Barr virus-transformed B cell lines. Preferential utilization of the most JH-proximal D segment (DQ52) and two unusual VH-related rearrangements.";
J. Exp. Med. 169:1391-1403(1989).
  Ensembl; ENSG00000130076; Homo sapiens. InterPro; IPR0077110; Ig-1ike. InterPro; IPR007979; Ig-c1. InterPro; IPR003906; Ig_MHC. InterPro; IPR003906; Ig_W. Pfam; PF07654; C1-8et; 4. SMART; SM00406; IGV; 1.
  PIR;
  PIR; C
   MEDLINE=93301610; PubMed=8315388; DOI=10.1084/jem.178.1.331; Hillson J.L., Karr N.S., Oppliger I.R., Mannik M., Sasso E.H. "The structural basis of germline-encoded VH3 immunoglobulin
   PubMed=2840480; DOI=10.1084/jem.168.1.229;
Bird J., Galili N., Link M., Stites D., Sklar J.;
"Continuing rearrangement but absence of somatic hypermutation
immunoglobulin genes of human B cell precursor leukemia.";
J. Exp. Med. 168:229-245(1988).
  Neale G.A., Kitchingman G.R.; "mRNA transcripts initiating within the human immunoglobulin the chain enhancer region contain a non-translatable exon and are extremely heterogeneous at the 5' end."; Nucleic Acids Res. 19:2427-2433(1991).
   "Restricted utilization of germ-line VH3 genes complementarity-determining regions (CDR3) in h immunoglobulin heavy chain rearrangements."; Eur. J. Immunol. 22:247-251(1992).
   PROSITE; PS50835; IG_LIKE; 5.
PROSITE; PS00290; IG_MHC; UNKNOWN
   Raaphorst F.M., Timmers Schuurman R.K.;
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  NUCLEOTIDE
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Exp. Med. 178:331-33(1993).

Exp. Med. AAH20240.1; -; mRNA.
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G36005; G36005.

PH1642; PH1643; PH1643.

PH1643; PH1644.

PH1646; PH1646.

PH1098; PH1646.

PH0098; PH1646.

PH0120; PH16120.

S15590; S15590.

S31116; S31116.

S31119; S31119.

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                         30
  DB
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   and short diverse
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DT 10-MAY
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DT 110-MAY
DT 110
RESULT AT
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ID QSMTV3
AC QSMTV3
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DT 01-FEB
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Best Local
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Q5M7V3;
Q5M7V3;
Q1-FEB-2005 (TrEMBLrel. 2
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MEDLINE-81013859; PubMed-6774332;
Lehman D.W., Putnam F.W.;

"Amino acid sequence of the variable region of a human mu chai
location of a possible JH segment.";

Proc. Natl. Acad. Sci. U.S.A. 77:3239-3243(1980).

-i- MISCELLANEOUS: This mu chain was isolated from the plasma
  21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence update)
10-MAY-2005 (Rel. 47, Last annotation update)
Ig heavy chain V-III region CAM.
Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Verte
   MOD_RES
NON_TER
SEQUENCE
   This Swiss-Prot entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation - the Buropean Bioinformatics Institute. There are no restrictions on its the Buropean Bioinformatics Institute. There are no restrictions on its use as long as its content is in no way modified and this statement is not
  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Buarchontoglires; Primates; Catarrhini; Hominidae;
  DOMAIN
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GO; GO:0003823; F:antigen binding; NAS.
GO; GO:0006952; F:lmmune response; NAS.
InterPro; IPR007110; Ig-11ke.
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SMART; SM00406; IGv; 1.
  NCBI_TaxID=9606;
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
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                               NUCLEOTIDE SEQUENCE
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SIMILARITY: Contains 1 Ig-like (immunoglobulin-like) domain.
  HUMAN
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  Similarity
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122 AA;
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RA Strausberg R.L., Peingold E.A., Grouse L.H., Derge J.G.,
RA Klauener R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
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RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Mhiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Bukesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
Butterfield Y.S.N., Krzywinski M.I., Skalska U., Smailus D.E.,
"Generation and initial analysis of more than 15,000 full-length human
and mouse cDNA sequences.";
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  Matches
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InterPro; IPR007110; Ig-like.
InterPro; IPR003597; Ig_cl.
InterPro; IPR003596; Ig_v.
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MEDLINE=98277139; PubMed=9614934; DOI=10.1006/clin.1998.4531;
MEDLINE=98277139; PubMed=9614934; DOI=10.1006/clin.1998.4531;
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01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Myosin-reactive immunoglobulin heavy chain variable
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09UL93;
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   Submitted (DEC-2004) to the EMBL/GenBank/DDBJ databases.

- FUNCTION: Beta-2-microglobulin is the beta-chain of major
- histocompatibility complex class I molecules (By similarity).

-1- SUBCELLULAR LOCATION: Secreted (By similarity).
   NIH MGC Project;
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   HUMAN
   Local
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   Natl. Acad. Sci.
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   116
  49
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  region
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A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

A Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

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A Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

A Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

A Raha S.S., Moßwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,

A Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

Vallalon D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,

A Villalon D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,

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Generation and initial analysis of more than 15,000 full-length human
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Best Local S
Matches 25
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PIR; PL0120; PL0120.
HSSP; P01772; 2FB4.
SMR; Q9UL93; 1-116.
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Interpro; IPR003596; Ig-v.
Interpro; IRR003596; Ig-v.
IRROGITE; PS50835; IG_IKE; 1.
  Bird J., Galili N., Link M., Stites D., Sklar J.;
"Continuing rearrangement but absence of somatic hypermutation immunoglobulin genes of human B cell precursor leukemia.";
J. Exp. Med. 168:229-245(1988).
EMBL; AF035021; AAD56257.1; -; mRNA.
   MRDLINE=93301610; PubMed=8315388; DOI=10.1084/jem.178.1.331; Hillson J.L., Karr N.S., Oppliger I.R., Mannik M., Sasso E.H.; Hillson J.L., Karr N.S. germline-encoded VH3 immunoglobulin binding to staphylococcal protein A."; C3.Exp. Med. 178:331-336(1993).
   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;
Muroidea; Muridae; Murinae; Rattus.
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PubMed=2840480;
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MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
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   Rattus norvegicus (Rat).
   Name=LOC299354;
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   Q510J0;
10-MAY-2005
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  VQLVQSGGGVVHPGRSLKLSCAGSGFTFS 30
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  116
116 AA,
  Conservative
  PRELIMINARY;
  116
   83.8%;
86.2%;
  12434 MW;
   , Created)
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, Last ann
  2
  Score 129; DB 2;
Pred. No. 1.1e-09;
   0DA0348154DD6061 CRC64;
   Mismatches
   sequence update) annotation updat
   465
   ⋛
   Length 116,
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   Gaps
   human
   0
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RESULT 7

Q65ZC9 H

Q65ZC9 H

AC Q65ZC9 C

Q65ZC9 C

Q65ZC9 C

Q65ZC9 C

D7 25-OCT-2

D7 25-OCT-2

D8 Single-C

GN Name=scr

GN Name=scr

GN Namesla

OC Homo.

OC Mammalla

OC HOMO.

OC NCDI-Tax

RN [1]

RP NUCLBOTH

RC STRAIN-C

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RA INTERPRO

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DR SMART; 8

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   Query Match
Best Local &
Matches 25
   STRAIN=Clq/7;

MEDLINE=97362799; PubMed=9219263; DOI=10.1038/nbt0797-629;

MEDLINE=97362799; PubMed=9219263; DOI=10.1038/nbt0797-629;

M KONTERMANN R.B., Wing M.G., Winter G.;

"Complement recruitment using bispecific diabodies.";

I Nat. Biotechnol. 15:629-631(1997).

R EMBL; Y13056; CAA73499; 11.

R EMBL; Y13056; CAA73499; 12.

R InterPro; IPR00359; 13.

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R InterPro; IPR003596; 13.

R InterPro; IPR003596; 13.

R SMART; SM00406; IGy; 2.

R SMART; SM00406; IGy; 2.

R SMART; SM00406; IGy; 2.

R SMART; SM00406; IG; LIKE; 2.
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Matches 24
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InterPro; IPRO0310; Ig-like.
InterPro; IPR003197; Ig_cl.
InterPro; IPR003596; Ig_v.
Pfam; PP07654; Cl-set; 3.
SMART; SM00409; IG; 2.
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SMART; PS50835; IG_LIKE; 4.
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SEQUENCE 465 AA; 51699 MW; 97C4533A213A424F CRC64;
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TISSUE-Thymns;
TISSUE-Thymns;
NIH MGC Project;
Submitted (DEC-2004) to the EMBL/GenBank/DDBJ databases.
-!- FUNCTION: Beta-2-microglobulin is the beta-chain of major
-i- subcratibility complex class I mblecules (By similarity).
-!- SUBCELLULAR LOCATION: Secreted (By similarity).
EMBL; BC088271; AAH8871.1; -; mRNA.
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Homo sapiens (Human).
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Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
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  Proc. Natl. Acad. Sci. [2]
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  -
   -
  25,
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   Similarity
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   Conservative
  82.5%;
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   83.8%;
   U.S.A. 99:16899-18903(2002)
   ک
   Last sequence update)
Last annotation update)
  Created)
   4:
   Score 127; DB 2;
Pred. No. 4.2e-09;
2; Mismatches 3
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Pred. No. 4.2e-09;
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   Mismatches
  240
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  Gaps
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RESULT

8

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RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
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RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Brownstein M.J., Ugdin T.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Brownstein M.J., Ugdin T.B., Toshiyuki S., Carrinci P., Prange C.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Richards S., Morley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Willalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Wilting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Blakesley R.W., Touchman J.W., Green B.D., Dickson M.C.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Schnerch A., Schein J.B., Jones S.J.M., Marra M.A.;
RT "Generation and initial analysis of more than 15,000 full-length human and mouse cDNA sequences";
RT proc. Marl Acad Sci. U.S. A 99.16899-1690170002)
RESULT 9
OSEFES HUMAN
OSEFES;
OSEFES;
OSEFES;
OT 10-MAY-2005 (TremBlrel. 30,
DT 10-MAY-2005 (TremBlrel. 30,
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  Query Match
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   Matches
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InterPro; IPR003597; Ig cl.
InterPro; IPR00306; Ig wHC.
InterPro; IPR003396; Ig v.
Pfam; PF07654; Cl-set; 2.
SMART; SM00409; IG; 3.
SMART; SM00407; IGC1; 3.
SMART; SM00406; IGV; 1.
PROSITE; PS50835; IG LIKE; 3.
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SMR; Q5BK12; 20-241.
GO; GO:0003823; F:antigen binding; IBA.
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Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurogna
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10-MAY-2005
   Q5BK12_RAT
Q5BK12;
  NCBI_TaxID=10116;
  Igha protein.
  NUCLEOTIDE SEQUENCE.
   rissu=spleen;
   RAT
   20
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  Similarity
   Project;
   PRELIMINARY
  Acad.
   Conservative
  82.5%;
  U.S.A. 99:16899-16903(2002).
   Last sequence update)
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   Created)
Last seq
  EMBL/GenBank/DDBJ databases.
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   Score 127; DB
Pred. No. 8.2e
3; Mismatches
 Created)
Last sequence update)
  PRT;
   8B146164AE8437C5
  PRT;
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  475
  3;
  8
   BB
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  update)
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  Rodentia; Sciurognathi;
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   0
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HY3J HUMAN
STANDAKU;
HY3J HUMAN
STANDAKU;
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21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 04, Last sequence update)
10-MAY-2005 (Rel. 47, Last annotation update)
10-MAY-2005 (Rel. 47, Last annotation update)
11 heavy chain V-III region HIL.
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Pfam; PF07686; V-set; 1.
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  Gaucher C., Klean
"commence determination
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Eukaryota; Metazoa; Chordata; Craniata;
Mammalia; Eutheria; Euarchontoglires; Pr
  PROTEIN SEQUENCE.

MEDILINE=79124695; PubMed=420800;

Chiu Y.-Y.H., Lopez de Castro J.A., Poljak R.J.;

"Amino acid sequence of the VH region of human myeloma
cryoimmunoglobulin IgG Hil.";

Biochemistry 18:553-560(1979).

-i- MISCELLANEOUS: This chain was isolated from an IgG1 myeloma
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  "Sequence determ
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  10-MAY-2005 (TrEMBLrel. 30, Last annotation update)
Anti-RhD monoclonal T125 gammal heavy chain precursor.
   This Swiss-Prot entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation - the European Bioinformatics Institute. There are no restrictions on its
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InterPro; IPR003006; Ig_MHC.
InterPro; IPR003596; Ig_v.
InterPro; IPR003596; Ig_v.
   InterPro; IPR003599; InterPro; IPR007110;
   removed.
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  as long as its content is
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   20
  1 QVQLVQSGGGVVHPGRSLKLSCAGSGFTF 29
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PS00290; IG_MHC; UNKNOWN_2.
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   1
20
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475
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Pred. No. 1.1e
2; Mismatches
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chain.
; 1367D400DC7D2859 CRC64;
   Potential.
  in no way
   Primates; Catarrhini; Hominidae;
  48
  1.1e-08
  Vertebrata; Euteleostomi;
  modified and
  DB 2;
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  Length 475;
  Indels
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  Query Match
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Matches 23
   Query Match
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Matches 24
   R SMR; P01771; 2-121.

R GO; GO:0005576; C:extracellular region; NAS.

R GO; GO:0003823; F:antigen binding; NAS.

R GO; GO:0003823; F:antigen binding; NAS.

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R InterPro; IPR007110; Ig-1ike.

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PROSITE; PS50835; IG LIKE; 1.

Direct protein sequencing; Immunoglobulin domain;

Immunoglobulin V region; Pyrrolidone carboxylic acid.

DOMAIN

1 112

Pyrrolidone carboxylic acid.

Pyrrolidone carboxylic acid.
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MEDLINE=74175307; PubMed=4208843;

Plorent G., Lehman D., Putnam F.W.;

"The switch point in mu heavy chains of human IgM immunoglobulins.";

Biochemistry 13:2482-2498(1974).
  HUMAN
   MOD_RES
   This Swiss-Prot entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation - the European Bioinformatics Institute. There are no restrictions on its
   21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence up
10-MAY-2005 (Rel. 47, Last annotation
1g heavy chain V-III region GA.
Homo sapiens (Human)
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SMR; P01769; 5-122.

SG); GO:0005576; C:extracellular region; GO:0003823; F:antigen binding; NAS.

GO; GO:0003823; F:tmmune response; NAS.

GO; GO:0006555; P:tmmune response; NAS.

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InterPro; IPR003596; Ig-v.
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  SEQÜENCE
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macroglobulin.

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  Eukaryota; Metazoa;
  NCBI_TaxID=9606;
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                            81.2%;
Score 125; DB 1;
Pred. No. 4.1e-09;
4; Mismatches ?
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Pred. No. 4
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Pyrrolidone carboxylic acid.
   PRT;
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  update)
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   carboxylic acid
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  Length 122;
   Length 121;
  Indels
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     0
   Gaps
     Gaps
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   0
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  InterPro; lrav; InterPro; lrav; I.
InterPro; lrav; I.
InterPro; lrav; I.
SMART; SM00406; IG_LIKB; 1.
SPOSTITE; PS50835; IG_LIKB; 1.
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Tmmunoglobulin V region; Pyrrolidone carboxylic acid.
Tmmunoglobulin V region; Pyrrolidone carboxylic acid.
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  GO; GO:0005576; C:extracellular region; GO; GO:0003823; F:antigen binding; NAS. GO; GO:0006955; P:immune response; NAS. InterPro; IPR007110; Ig-like. InterPro; IPR003596; Ig-v. SMART; SM00406; IGv; 1.
  This Swiss-Prot entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the El the European Bioinformatics Institute. There are no resti
   and 1.0-A resolution.";
J. Mol. Biol. 141:369-391(1980).
-!- SIMILARITY: Contains 1 Ig-like (immunoglobulin-like) domain.
  P01772;
21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence update)
10-MAY-2005 (Rel. 47, Last annotation updat
Ig heavy chain V-III region KOL.
Homo sapiens (Human).
TURN
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   Marquart M., Deisenhofer J., Huber R., Palm W.; "Crystallographic refinement and atomic models of the intact immunoglobulin molecule Kol and its antigen-binding fragment
   Schmidt W.E., Jung H.-D., Palm W., Hilschmann N.;
"Three-dimensional structure determination of antibodies. Printered in the structure of crystallized monoclonal immunoglobulin IgG1 KOL, hoppe-Seyler's Z. Physiol. Chem. 364:713-747(1983).
  use as
  MEDLINE=81072295; PubMed=7441755;
  X-RAY CRYSTALLOGRAPHY (1.9 ANGSTROMS).
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  A02055; G1HUKL
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||:||:||| | ||:||:|| ||:||
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  28
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   STANDARD;
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  110
126
126
126
127
127
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127
139
142
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173
  Chordata; Craniata; Vertebrata; Euteleostomi; Euarchontoglires; Primates; Catarrhini; Hominidae;
   BONDS
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  update)
  NAS.
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  Immunoglobulin domain;
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  the EMBL outstation
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InterPro; IPR003599; Ig.
InterPro; IPR003599; Ig.-like.
InterPro; IPR003597; Ig.-like.
InterPro; IPR003597; Ig.-di.
InterPro; IPR003596; Ig_MHC.
InterPro; IPR003596; Ig_v.
Pfam; PF07654; C1-set; Z.
SMART; SM004409; IG; 3.
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HELIX
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   Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;
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  NIH MGC Project
  STRAIN=FVB/N;
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   24;
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78
88
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  TISSUE=Colon;
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90
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108
116
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  13718 MW; E4D71B52B16F8776 CRC64;
   81.2%;
80.0%;
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Last annotation updat
   ω
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   Mismatches
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   update)
  Length 126;
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Matches 24
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SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG LIKE; 4.
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203
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NUCLEOTIDE SEQUENCE.

MEDLINE=98277139; PubMed=9614934; DOI=10.1006/clin.1998.4531;

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   InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 1.
   Bur. J. Immunol. 22:247-251(1992).
EMBL; AP035024; AAD56260.1; -; mRNA
PIR; S78486; S78486.
   complementarity-determining regions (CDR3) in human feral a lumanimmunoglobulin heavy chair
   Raaphorst F.M., Timmers E., Kenter M.J., Schuurman R.K.;
  Wu X., Liu B., Young D.C.;
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  retus.";
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   immunoglobulin heavy chain rearrangements.";
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  21
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PRELIMINARY;
(TrEMBLrel. 31,
(TrEMBLrel. 31,

Last sequence update)

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  RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
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RA Robak S.A., WcEwan P.J., McKernan K.J., Malek J.A., Gunarathe P.H.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
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Copyright (c) 1993 - 2005 Compugen Ltd.
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| Query M   | NESULT 2 US-09-534-717- Sequence 66: Patent No. (6) PAPLICANT: FILE REPERI CURRENT APICURENT FILE REPERI CURRENT FILE ARLIER FILE ARLIER FILE ARLIER FILE SOFTWARE: (1) SEQ ID NO 6: SEQ ID NO 6: SEQ ID NO 6: TYPE: PRT ORGANISM: US-09-534-717                                                                                                                                                              | Query Mat<br>Best Loca<br>Matches<br>Qy                                                                       | RESULT 1  US-09-534-717-617  US-09-534-717-617  ; Sequence 617, Application US/0 ; Patent No. 6914128 ; Patent No. 6914128 ; GENERAL INFORMATION: APPLICANT: JOChen, Salfeld et ; TITLE OF INVENTION: Human Ant ; FILE REFERNCE: BB1-093CP ; CURRENT PILING DATE: 2000-03 ; CURRENT FILING DATE: 2000-03 ; EARLIER APPLICATION NUMBER: U PARLIER OF SEQ ID NOS: 675 ; NUMBER OF SEQ ID NOS: 675 ; SOFTMARE: PATENTIN Ver. 2.0 ; SEQ ID NO 617 ; LENGTH: 97 ; TYPE: PRT ; ORGANISM: Homo sapiens US-09-534-717-617                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                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| Match     | SULT 2 Sequence 667, Application Sequence 667, Application 10. 6914128 GENERAL INFORMATION: APPLICANT: Jochen, FITLE OF INVENTION: FILE REFERENCE: BBI CURRENT APPLICATION CURRENT FILING DATE EARLIER FILING DATE EARLIER FILING DATE EARLIER FILING DATE EARLIER FILING DATE EARLIER FILING DATE EARLIER FILING DATE OF SEQ ID NO 667 SEQ ID NO 667 LENGTH: 98 TYPE: PRT ORGANISM: Homo sap 199-534-717-667 | ch<br>1 s<br>26                                                                                               | TT 1 9-534-717-617 9-534-717-617 9-534-717-617 9-100-600 9-100-600 9-100-600 9-100-600 9-100-600 9-100-600 9-100-600 9-100-600 9-100-600 9-100-600 9-100-600 9-100-600 9-100-600 9-100-600 9-100-600 9-100-600 9-100-600 9-100-600 9-100-600 9-100-600 9-100-600 9-100-600 9-100-600 9-100-600 9-100-600 9-100-600 9-100-600 9-100-600 9-100-600 9-100-600 9-100-600 9-100-600 9-100-600 9-100-600 9-100-600 9-100-600 9-100-600 9-100-600 9-100-600 9-100-600 9-100-600 9-100-600 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|           | e e z zoza t                                                                                                                                                                                                                                                                                                                                                                                                  | h 89.6%; Score 138; Similarity 86.7%; Pred. No. 26; Conservative 3; Mismatcl 1 QVQLVQSGGGVVHPGRSLKLSCAGSGFTFS | 14-717-617 10ce 617, Application US/09534717 12 NO. 6914128 12 NO. 6914128 13 OF INVENTION: Salfeld et al. 13 OF INVENTION: Human Antibodies 13 OF INVENTION: Human Antibodies 13 OF INVENTION HUMBER: US/09/53 13 NT APPLICATION NUMBER: US/09/53 13 NT FILING DATE: 2000-03-46 1ER APPLICATION NUMBER: 60/126,6 1ER FILING DATE: March 25, 1999 1ER OF SEQ ID NOS: 675 1ER OF SEQ ID NOS: 675 1ER OF SEQ ID NOS: 675 1ER OF SEQ ID NOS: 675 1ER OF SEQ ID NOS: 675 1ER OF SEQ ID NOS: 675 1ER OF SEQ ID NOS: 675 1ER OF SEQ ID NOS: 675 1ER OF SEQ ID NOS: 675 1ER OF SEQ ID NOS: 675 1ER OF SEQ ID NOS: 675 1ER OF SEQ ID NOS: 675 1ER OF SEQ ID NOS: 675 1ER OF SEQ ID NOS: 675 1ER OF SEQ ID NOS: 675 1ER OF SEQ ID NOS: 675 1ER OF SEQ ID NOS: 675 1ER OF SEQ ID NOS: 675 1ER OF SEQ ID NOS: 675 1ER OF SEQ ID NOS: 675 1ER OF SEQ ID NOS: 675 1ER OF SEQ ID NOS: 675 1ER OF SEQ ID NOS: 675 1ER OF SEQ ID NOS: 675 1ER OF SEQ ID NOS: 675 1ER OF SEQ ID NOS: 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68<br>1-100<br>1-100<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-106<br>1-1 |
| Length 98 | IL-12                                                                                                                                                                                                                                                                                                                                                                                                         | Length 97;<br>Indels                                                                                          | IL-12                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      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| ••        | And Methods                                                                                                                                                                                                                                                                                                                                                                                                   | 0,                                                                                                            | And Methods                                                                                                                                                                                                                                                                                                                                                                                                                                                                                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|           | nods For                                                                                                                                                                                                                                                                                                                                                                                                      | Gaps                                                                                                          | ods For                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        |            | 68, Appl 94, Appl 100, App 100, App 62, Appl 68, Appl 68, Appl 106, App 106, App 106, App 116, Appl 153, Appl 153, Appl 118, Appl 118, Appl 124, Appl 25, Appl 25, Appl 26, Appl                                                                                                                                                                                                                                               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|           | Producing                                                                                                                                                                                                                                                                                                                                                                                                     | 0,                                                                                                            | Producing                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  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|           | Ing                                                                                                                                                                                                                                                                                                                                                                                                           |                                                                                                               | 'ng                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        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Best Local Similarity

Conservative

90.0%;

Pred. No. 2.5e-11; 1, Mismatches 2;

Indels

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Gaps

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  PILE REFERENCE: 213839-00013
CURRENT APPLICATION NUMBER: US/09/726,219A
CURRENT FILING DATE: 2000-11-28
PRIOR APPLICATION NUMBER: GB 9015198.6
PRIOR FILING DATE: 1990-07-10
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR FILING DATE: 1990-10-19
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR FILING DATE: 1990-10-19
PRIOR APPLICATION NUMBER: GB 9024503.6
PRIOR APPLICATION NUMBER: GB 9024503.6
PRIOR APPLICATION NUMBER: GB 9024744.9
PRIOR APPLICATION NUMBER: GB 9104744.9
PRIOR FILING DATE: 1991-03-06
  RESULT 3
US-09-726-219A-167
  US-09-534-717-23
GENERAL INFORMATION:
APPLICANT: Jochen, Salfeld et al.
   Sequence 23, Application US/09534717 Patent No. 6914128
   Sequence 167, Application US/09726219A Patent No. 6806079
   SOPTWARE: PatentIn version 3.1
SEQ ID NO 167
LENGTH: 115
  GENERAL INFORMATION:
   Matches
  Query Match
   TYPE: PRT
ORGANISM: Homo sapiens
-09-726-219A-167
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APPLICANT:
APPLICANT:
   APPLICANT:
APPLICANT:
APPLICANT:
   PRIOR APPLICATION NUMBER: US 08/484,893
PRIOR FILING DATE: 1995-06-07
NUMBER OF SEQ ID NOS: 272
  PRIOR APPLICATION NUMBER: GB 9110549.4
PRIOR FILING DATE: 1991-05-15
PRIOR APPLICATION NUMBER: PCT/GB91/01134
PRIOR FILING DATE: 1991-07-10
PRIOR APPLICATION NUMBER: US 07/971,857
PRIOR FILING DATE: 1993-01-08
PRIOR FILING DATE: 1993-01-08
   APPLICANT: Winter, Gregory
APPLICANT: Bonert, Timothy
TITLE OF INVENTION: Methods for Producing Members of Specific Binding Pairs
FILE REFERENCE: 213839-00013
   APPLICANT:
   APPLICANT:
   APPLICANT: Cambridge Antibody Technology Limited
   PPLICANT:
   Local
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  1 QVQLVQSGGGVVHPGRSLKLSCAGSGFTFS 30
  Similarity
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   Clackson, Timothy Chiswell, David
   Jackson, Ronald
  Griffiths, Andrew
   Johnson, Kevin
   Pope, Anthony
  McCafferty, John
   Holliger, Kasper
   Conservative
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   1; Mismatches
  Score 137; DB 2;
Pred. No. 2.9e-11;
   30
  Length 115;
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; ORGANISM: Homo sapiens
US-09-534-717-33
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                                    US-09-534-717-37
   US-09-534-717-37
  US-09-534-717-33
  ; ORGANISM: Homo sapiens
US-09-534-717-23
  CURRENT APPLICATION NUMBER: US/09/534,717
CURRENT FILING DATE: 2000-03-24
EARLIER APPLICATION NUMBER: 60/126,603
EARLIER FILING DATE: March 25, 1999
NUMBER OF SEQ ID NOS: 675
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 37
  CURRENT APPLICATION NUMBER: US/09/534,717
CURRENT FILING DATE: 2000-03-24
EARLIER APPLICATION NUMBER: 60/126,603
EARLIER FILING DATE: March 25, 1999
NUMBER OF SEQ ID NOS: 675
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 33
  TITLE OF INVENTION: Human Antibodies That Bind Human IL-12 And Methods For Producing FILE REFERENCE: BBI-093CP CURRENT APPLICATION NUMBER: US/09/534,717 CURRENT FILING DATE: 2000-03-24 EARLIER APPLICATION NUMBER: 60/126,603 EARLIER FILING DATE: March 25, 1999 NUMBER OF SEQ ID NOS: 675 SOFTWARE: Patentin Ver. 2.0 SEQ ID NO 23 LENGTH: 115
   Sequence 33, Application US/09534717
Patent No. 6914128
GENERAL INFORMATION:
   Sequence 37, App
Patent No. 6914
  GENERAL INFORMATION:
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   Query Match
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  Matches
   Query Match
Best Local Similarity
Query Match
   APPLICANT: Jochen, Salfeld et al.
TITLE OF INVENTION: Human Antibodies That Bind Human IL-12 And Methods For Producing
FILE REFERENCE: BBI-093CP
   APPLICANT: Jochen, Salfeld et al.
TITLE OF INVENTION: Human Antibodies That
FILE REFERENCE: BBI-093CP
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ORGANISM: Homo sapiens
  TYPE: PRT
   LENGTH:
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  Similarity 90.0
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  Application US/09534717
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 Score 137;
   Score 137; DB 2;
Pred. No. 2.9e-11;
   Score 137; DB 2;
Pred. No. 2.9e-11;
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Best Local Matches :

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90.0%;

Pred. No. 2.90 1; Mismatches

No. 2.9e-11; smatches 2;

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QVQLVQSGGGVVHPGRSLKLSCAGSGFTFS 30

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US-09-534-717-39
   RESULT 9
US-09-534-717-43
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  US-09-534-717-41
   PATENT NO. 6914128
GENERAL INFORMATION: Salfeld et al.
APPLICANT: Jochen, Salfeld et al.
TITLE OF INVENTION: Human Antibodies That Bind Human IL-12 And Methods For Producing PILE REFERENCE: BBI-093CP
CURRENT FILING NUMBER: US/09/534,717
CURRENT FILING DATE: MONO-03-24
EARLIER APPLICATION NUMBER: 60/126,603
EARLIER FILING DATE: March 25, 1999
NUMBER OF SEQ ID NOS: 675
SOCTWARE: PatentIn Ver. 2.0
SEQ ID NO 39
LENGTH: 115
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SOFTWARE: PATENTIN VEr. 2.0
SEQ ID NO 41
LENGTH: 115
  Patent No. 6914128
GENERAL INFORMATION:
                               Sequence 43, Application US/09534717
Patent No. 6914128
GENERAL INFORMATION:
  Sequence 41, Application US/09534717
   Query Match
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   Query Match
   TYPE: PRT
ORGANISM: Homo sapiens
-09-534-717-41
   APPLICANT: Jochen, Salfeld et al.
TITLE OF INVENTION: Human Antibodies That Bind Human IL-12
TITLE OF INVENTION: Human Antibodies That Bind Human IL-12
TILE REFERENCE: BBI-093CP
CURRENT APPLICATION NUMBER: US/09/534,717
CURRENT FILING DATE: 2000-03-24
EARLIER APPLICATION NUMBER: 60/126,603
EARLIER FILING DATE: March 25, 1999
  ORGANISM: Homo sapiens
APPLICANT: Jochen, Salfeld et al.
TITLE OF INVENTION: Human Antibodies That Bind Human IL-12 And Methods For Producing
  TYPE: PRT
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27; Conserv
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   Application US/09534717
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Pred. No. 2.9e-11;
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Pred. No. 2.9e-11;
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   Length 115;
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   Gaps
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   0;
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APPLICANT: JOCHEN, Salfeld et al.
TITLE OF INVENTION: Human Antibodies That Bind Hum
FILE REFERENCE: BBI-093CP
CURRENT APPLICATION NUMBER: US/09/534,717
CURRENT FILING DATE: 2000-03-24
EARLIER APPLICATION NUMBER: 60/126,603
EARLIER FILING DATE: March 25, 1999
NUMBER OF EGG ID NOS: 675
SOFTWARE: PatentIn Ver. 2.0
1 SEQ ID NO 47
   RESULT 10
US-09-534-717-45
; Sequence 45, A
  LENGTH: 115; TYPE: PRT; ORGANISM: Homo sapiens US-09-534-717-47
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  8
   ; ORGANISM: Homo sapiens US-09-534-717-43
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   ; ORGANISM: Homo sapiens
US-09-534-717-45
  US-09-534-717-47
  SEQ ID NO 43
LENGTH: 115
  GENERAL INFORMATION:
   SEQ ID NO 45
  Patent No. 6914128
GENERAL INFORMATION:
   Matches
  Query Match
Best Local
  Matches
  APPLICANT: Jochen, Salfeld et al.

APPLICANT: Jochen, Salfeld et al.

TITLE OF INVENTION: Human Antibodies That Bind Human IL-12 And Methods For Producing FILE REFERENCE: BBI-093CP
CURRENT APPLICATION NUMBER: US/09/534,717
CURRENT FILING DATE: 2000-03-24
EARLIER APPLICATION NUMBER: 60/126,603
EARLIER FILING DATE: March 25, 1999
NUMBER OF SEQ ID NOS: 675
SOFTWARE: Patentin Ver. 2.0
Query Match
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   Sequence 47, Application US/09534717 Patent No. 6914128
  Query Match
   FILE REFERENCE; BBI-093CP
CURRENT APPLICATION NUMBER: US/09/534,717
CURRENT FILING DATE: 2000-03-24
EARLIER APPLICATION NUMBER: 60/126,603
EARLIER FILING DATE: March 25, 1999
NUMBER OF SEQ.-ID NOS: 675
SOFTWARE: Patentin Ver. 2.0
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1 Similarity 90.0%;
27; Conservation
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  27; Conservative
  similarity 90.0%;
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  QVQLVQSGGGVVQPGRSLRLSCAASGFTFS 30
   Application US/09534717
 89.0%;
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Pred. No. 2.9e-11;
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Pred. No. 2.9e-11;
 Score 137; DB 2;
Pred. No. 2.9e-11;
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Sequence 53, Application US/09534717

Patent No. 6914128

GENERAL INFORMATION:
APPLICANT: Jochen, Salfeld et al.
ITILS OF INVENTION: Human Antibodies That Bind Human IL-
PILE REFERENCE: BBI-093CP
CURRENT APPLICATION NUMBER: US/09/534,717

CURRENT FILING DATE: 2000-03-24

EARLIER APPLICATION NUMBER: 60/126,603

EARLIER APPLICATION NUMBER: 50/126,603

EARLIER APPLICATION NUMBER: 50/126,603

SEQID NO 53

SOFTWARE: PATENTIN DATE: March 25, 199

NUMBER OF SEQ ID NOS: 675

SOFTWARE: PATENTIN Ver. 2.0
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   US-09-534-717-55
  RESULT 14
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  US-09-534-717-53
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US-09-534-717-51
  US-09-534-717-51
  RESULT 12
Sequence 55, Application US/09534717
Patent No. 6914128
GENERAL INFORMATION:
APPLICANT: Jochen, Salfeld et al.
TITLE OF INVENTION: Human Antibodies That Bind Human
FILE REFERENCE: BBI-093CP
   Query Match
Best Local Similarity
Matches 27; Conserv
  SOFTWARE: P
   GENERAL INFORMATION:
   Sequence 51, Appli
Patent No. 6914128
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   Matches
   Query Match
   CURRENT APPLICATION NUMBER: US/09/534,717
CURRENT FILING DATE: 2000-03-24
EARLIER APPLICATION NUMBER: 60/126,603
EARLIER FILING DATE: March 25, 1999
NUMBER OF SEQ ID NOS: 675
SOFTMARE: Patentin Ver. 2.0
   APPLICANT: Jochen, Salfeld et al.
TITLE OF INVENTION: Human Antibodies That Bind
FILE REFERENCE: BBI-093CP
   TYPE: PRT
  TYPE: PRT
  LENGTH:
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l Similarity 90.0%;
27; Conservative
   27;
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   QVQLVQSGGGVVQPGRSLRLSCAASGFTFS 30
  Application US/09534717
   Conservative
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   90.0%;
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   1,
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   DB 2;
2.9e-11;
  Human IL-12 And Methods For Producing
   ν
~
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                   IL-12
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                   And Methods For Producing
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   0
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   Gaps
   Gaps
   Gaps
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   0
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; ORGANISM: Homo sapiens
US-09-534-717-57
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Search completed: November 21, Job time: 11.3187 secs
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  GENERAL INFORMATION:
  APPLICANT: Jochen, Salfeld et al.
TITLE OF INVENTION: Human Antibodies That Bind Human IL-12 And
FILE REFERENCE BBI-093CP
CURRENT APPLICATION NUMBER: US/09/534,717
CURRENT FILING DATE: 2000-03-24
EARLIER APPLICATION NUMBER: 60/126,603
EARLIER FILING DATE: March 25, 1999
NUMBER OF SEQ ID NOS: 675
  Sequence 57, Appl Patent No. 691412
  CURRENT APPLICATION NUMBER: US/09/534,717
CURRENT FILING DATE: 2000-03-24
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EARLIER FILING DATE: March 25, 1999
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US-10-884-830-617
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  Application US/10632706
DS20040175385A1
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| 137                                                                                                                                                                                                                                                                                                                                                                                                                                                     |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       | <b>4</b> 5 | 44    | 43   | 42   | 41   | 40   | 9    | 38   | 37   | 36   | 35          | 3.4<br>4. | ü          | 32         | 31          | 30          | 29          | 20         |
|---------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|---------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|------------|-------|------|------|------|------|------|------|------|------|-------------|-----------|------------|------------|-------------|-------------|-------------|------------|
| 9.0 115 4 US-10-803-622<br>9.0 115 5 US-10-804-83<br>9.0 115 5 US-10-804-83 | 9.0 115 4 US-10-803-622-167 Sequence 9.0 115 4 US-10-803-653-167 Sequence 9.0 115 5 US-10-804-830-23 Sequence 9.0 115 5 US-10-804-830-33 Sequence 9.0 115 5 US-10-804-830-37 Sequence 9.0 115 5 US-10-804-830-37 Sequence 9.0 115 5 US-10-804-830-41 Sequence 9.0 115 5 US-10-804-830-43 Sequence 9.0 115 5 US-10-804-830-45 Sequence 9.0 115 5 US-10-804-830-57 Sequence 9.0 115 5 US-10-804-830-57 Sequence 9.0 115 5 US-10-804-830-59 Sequence 9.0 115 5 US-10-804-830-59 Sequence 9.0 115 5 US-10-804-830-63 Sequence 9.0 115 5 US-10-804-830-63 Sequence 9.0 115 5 US-10-804-830-63 Sequence 9.0 115 5 US-10-804-830-63 Sequence 9.0 115 5 US-10-804-830-63 Sequence 9.0 115 5 US-10-804-830-63 Sequence 9.0 115 5 US-10-804-830-63 Sequence                                     | 137        | 137   | 137  | 137  | 137  | 137  | 137  | 137  | 137  | 137  | 137         | 137       | 137        | 137        | 137         | 137         | 137         | 101        |
| 4 US-10-803-623 4 US-10-803-623 5 US-10-884-833                                                                                                                                                                         | 4 US-10-803-622-167 Sequence 4 US-10-803-623-167 Sequence 5 US-10-884-830-23 Sequence 5 US-10-884-830-37 Sequence 5 US-10-884-830-37 Sequence 6 US-10-884-830-41 Sequence 7 US-10-884-830-41 Sequence 7 US-10-884-830-45 Sequence 8 US-10-884-830-47 Sequence 8 US-10-884-830-57 Sequence 9 US-10-884-830-51 Sequence 1 US-10-884-830-52 Sequence 1 US-10-884-830-53 Sequence 1 US-10-884-830-53 Sequence 1 US-10-884-830-59 Sequence 1 US-10-884-830-59 Sequence 1 US-10-884-830-59 Sequence 1 US-10-884-830-59 Sequence 1 US-10-884-830-59 Sequence 1 US-10-884-830-59 Sequence 1 US-10-884-830-59 Sequence 1 US-10-884-830-59 Sequence 1 US-10-884-830-63 Sequence 1 US-10-884-830-63 Sequence 1 US-10-884-830-63 Sequence 1 US-10-884-830-63 Sequence 1 US-10-884-830-63 Sequence | 89.0       | 89.0  | 89.0 | 89.0 | 89.0 | 89.0 | 89.0 | 89.0 | 89.0 | 89.0 | 89.0        | 89.0      | 89.0       | 89.0       | 89.0        | 89.0        | 89.0        | 07.0       |
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|                                                                                                                                                                                                                                                                                                                                                                                                                                                         |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       |            |       |      |      |      |      |      |      |      |      | -10-884-830 |           | -10-884-83 | -10-884-83 | -10-884-830 | -10-803-653 | -10-803-622 | 10 001 000 |
| 167, App 23, App 23, App 23, App 33, App 34, App 41, App 41, App 45, App 47, App 47, App 51, App 55, App 55, App 55, App 56, App 66, App                                                                                                                                                                                                                                                                                                                |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       | Appl       | Today | Tddy | Appı | Tddv | Tddv | Appl | Appl | Tddv | Iddy | Tddv        | rddv      | Tddv       | Tddv       | Tddy        | ddv         | App         |            |

#### ALIGNMENTS

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OTHER INFORMATION: single chain antibody fragment US-10-632-706-158
  Sequence 162, Application, Publication No. US20040175, GENERAL INFORMATION:
APPLICANT: MARKS, JAMES I APPLICANT: AMERSDORFER,
   GENERAL INFORMATION:

APPLICANT: MARKS, JAMES D.

APPLICANT: MARKS, JAMES D.

APPLICANT: MARKS, JAMES D.

APPLICANT: MARKS, JAMES D.

TITLE OF INVENTION: THERAPEUTIC MONOCLONAL ANTIBODIES THAT NEUTRALIZE BOTULINUM
TITLE OF INVENTION: NEUROTOXINS
TILE REFERENCE: 4077-895120US
CURRENT APPLICATION NUMBER: US/10/632,706
CURRENT APPLICATION NUMBER: US 60/400,721
PRIOR APPLICATION NUMBER: US 60/400,721
PRIOR APPLICATION NUMBER: US 60/400,721
PRIOR APPLICATION NUMBER: US 60/400,721
PRIOR FILING DATE: 2002-08-01
PRIOR FILING DATE: 1998-08-31
NUMBER OF SEQ ID NOS: 278

PRIOR PRICE TELEVILLE 1998-08-31
NUMBER OF SEQ ID NOS: 278
   SOFTWARE: PatentIn version 3.2
SEQ ID NO 158
LENGTH: 30
APPLICANT: AMERSDORFER, PETER
TITLE OF INVENTION: THERAPEUTIC MONOCLONAL ANTIBODIES THAT NEUTRALIZE BOTULINUM
TITLE OF INVENTION: NEUROTOXINS
FILE REFERENCE: 407T-895120US
CURRENT APPLICATION NUMBER: US/10/632,706
CURRENT FILING DATE: 2003-08-01
PRIOR APPLICATION NUMBER: US 60/400,721
PRIOR PLICATION NUMBER: US 09/144,806
  ORGANISM: Artificial FEATURE:
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TITLE OF INVENTION: THERAPEUTIC MONOCLONAL ANTIBODIES THAT NEUTRALIZE BOTULINUM
TITLE OF INVENTION: NEUROTOXINS
FILE REFERENCE: 4077-895120US
CURRENT APPLICATION NUMBER: US/10/632,706
CURRENT FILING DATE: 2003-08-01
PRIOR APPLICATION NUMBER: US 60/400,721
PRIOR FILING DATE: 2002-08-01
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PRIOR FILING DATE: 2002-08-01
PRIOR FILING DATE: 1908-01
NUMBER: SOFTWARE: Patentin version 3.2
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LENGTH: 30
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Sequence 170, Application US/10632706
Publication No. US20040175385A1
GENERAL INFORMATION:
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US-10-632-706-166
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   Sequence 166, Application US/10632706 Publication No. US20040175385A1
   PRIOR FILING DATE: 1998-08-31
NUMBER OF SEQ ID NOS: 278
SOFTWARE: PatentIn version 3.2
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APPLICANT: MARKS, JAMES D.
APPLICANT: MARKS, JAMES D.
APPLICANT: AMERSDORFER, PETER
TITLE OF INVENTION: THERAPEUTIC MONOCLONAL ANTIBODIES THAT NEUTRALIZE BOTULINUM
TITLE OF INVENTION: NEUROTOXXINS
FILE REFERENCE: 407T-895120US
CURRENT APPLICATION NUMBER: US/10/632,706
CURRENT FILING DATE: 2003-08-01
PRIOR APPLICATION NUMBER: US 60/400,721
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NUMBER OF SEQ ID NOS: 278
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FILE REFERENCE: MJI-002

CURRENT APPLICATION NUMBER: US/10/911,838

CURRENT FILING DATE: 2004-08-04

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PRIOR FILING DATE: 2004-04-26

PRIOR APPLICATION NUMBER: US 60/545670

PRIOR FILING DATE: 2004-02-18

PRIOR FILING DATE: 2004-02-19

PRIOR FILING DATE: 2003-10-09

PRIOR FILING DATE: 2003-10-09
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Publication No. US20050004354A1
GENERAL INFORMATION:
APPLICANT: Jochen, Salfeld et al.
TITLE REFERENCE: BBI-093CP
FILE REFERENCE: BBI-093CP
CURRENT APPLICATION NUMBER: US/10/884,830
CURRENT FILING DATE: 2004-07-01
PRIOR FILING DATE: 2004-07-01
PRIOR PRICATION NUMBER: US/09/534,717
PRIOR FILING DATE: 2000-03-24
PRIOR APPLICATION NUMBER: 60/126,603
PRIOR PRILING DATE: March 25, 1999
PRIOR FILING DATE: March 25, 1999
   SEQ ID NO 617
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LENGTH: 30
   Sequence 18, Application US/10911838 Publication No. US20050069869A1
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  APPLICANT:
APPLICANT:
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  APPLICANT: BABCOCK, Gregory J.
APPLICANT: THOMAS, JR., William D.
APPLICANT: SULLIVAN, John
TITLE OF INVENTION: SARS NUCLEIC ACIDS,
TITLE OF INVENTION: AND USES THEREOF
   NUMBER OF SEQ ID NOS: 675
SOFTWARE: PatentIn Ver. 2.0
  APPLICANT:
   APPLICANT: AMBROSINO, Donna
  ORGANISM: Homo sapiens
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   OTHER INFORMATION: single chain antibody fragment
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US-10-982-359-18
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; Publication No. US20050112694A1
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Publication No. US20050112694A1
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NUMBER OF SEQ ID NOS: 148
SOFTWARE: FastSEQ for Windows Version 4.0
   APPLICANT: Carter, Paul J.
APPLICANT: Zhou, Hongxing
TITLE OP INVENTION: ANTIBODIES THAT BIND INTERLEUKIN-4 RECEPTOR
FILE REFERENCE: 3492-A
FILE REFERENCE: 3492-A
CURRENT APPLICATION NUMBER: US/10/982,359
CURRENT FILING DATE: 2004-11-04
PRIOR APPLICATION NUMBER: 60/518,166
PRIOR PILING DATE: 2003-11-07
PRIOR FILING DATE: 2003-11-07
  CURRENT APPLICATION NUMBER: US/10/982,359
CURRENT FILING DATE: 2004-11-04
PRIOR APPLICATION NUMBER: 60/518,166
PRIOR FILING DATE: 2003-11-07
  APPLICANT: Carter, Paul J.
APPLICANT: Zhou, Hongxing
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  APPLICANT: Carter, Paul J.
APPLICANT: Zhou, Hongxing
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FILE REFERENCE: 3492-A
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CURRENT FILING DATE: 2004-11-04
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PRIOR FILING DATE: 2003-11-07
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APPLICANT: Zhou, Hongxing
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CURRENT FILING DATE: 2004-11-04
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PRIOR APPLICATION NUMBER: 60/518,166
PRIOR PILING DATE: 2003-11-07
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APPLICANT: Carter, Paul J.
APPLICANT: Zhou, Hongxing
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   CURRENT APPLICATION NUMBER: US/10/982,359
CURRENT FILING DATE: 2004-11-04
PRIOR APPLICATION NUMBER: 60/518,166
PRIOR FILING DATE: 2003-11-07
NUMBER OF SEQ ID NOS: 77
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APPLICANT: Zhou, Hongxing
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APPLICANT: Carter, Paul J.

APPLICANT: Zhou, Hongxing

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APPLICANT: Zhou, Hongxing
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| Jion US/11144 50244408A1 Fean Penelope E. James D. James D. AMTIBODIES T PP2 RUMBER: US/1 RUMBER: US/1                                                                                                                                               | 81.2%;<br>76.7%;<br>ive<br>HPGRSLK<br>    ;<br>QPGRSLR                  | tion US/107217<br>050249729A1<br>ER KABUSHIKL R<br>ANTI TRAIL-R<br>1573-PCT<br>NUMBER: US/10<br>2003-11-26<br>UMBER: JP2001-<br>0001-05-18<br>UMBER: JP2001-<br>0001-08-09<br>2001-08-09<br>2001-08-09<br>10MBER: JP2001-<br>2001-10-11<br>5:: 45<br>Ver. 2.1                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                |
| 4424<br>TO<br>711/                                                                                                                                                                                                                                   | Score<br>Pred.<br>5; Mi<br>LSCAGS<br>                                   | 0721763<br>11KI KAISHA<br>AIL-R ANTIBODY<br>105/10/721,763<br>12001-150213<br>1802001-243040<br>99<br>2001-314489<br>11                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    | ALIGNMENTS | US-10-997-201A-6 US-11-077-978-7 US-11-077-978-2 US-11-079-978-3 US-10-932-334-83 US-10-932-334-7 US-10-932-334-7 US-10-932-334-7 US-10-932-334-7 US-10-932-334-7 US-10-932-334-7 US-10-932-334-7 US-10-932-334-7 US-10-932-334-7 US-10-932-334-7 US-10-932-334-7 US-10-932-334-7 US-10-932-334-7                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             |
| 8<br>INSULIN-LIKE GF                                                                                                                                                                                                                                 | 125; DB 1;<br>No. 9e-09;<br>smatches 2<br>GPTFS 30<br>     <br>GPTFS 49 | :3 ×                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       | STK        | 2016-7<br>978-7<br>978-2<br>978-3<br>334-3<br>334-9<br>334-9<br>334-9<br>334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>1334-7<br>133 |
| GROWTH FACTOR                                                                                                                                                                                                                                        | Length 139                                                              |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       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| H                                                                                                                                                                                                                                                    | B 0;                                                                    |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       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                                                                                                                                                                                                                                                                                                                                     |
| RECEPTOR                                                                                                                                                                                                                                             | Gарв                                                                    |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       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Sequence 4, Application US/10789273

Publication No. US20050249725A1

GENERAL INFORMATION:

APPLICANT: Basi, Guriq

APPLICANT: Saldanha, Jose

APPLICANT: Yednock, Ted

TITLE OF INVENTION: HUMANIZED ANTIBODIES THAT RECOGNIZE

TITLE OF INVENTION: BETA-AMYLOID PEPTIDE

FILE REFERENCE: ELM-002CP

CURRENT FILING DATE: 2004-02-27

PRIOR APPLICATION NUMBER: US/10/388,389

PRIOR PILING DATE: 2003-03-12

PRIOR PILING DATE: 2003-03-12
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SEQ ID NO 50
LENGTH: 473
TYPE: PRT
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US-11-144-248-50
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US-11-144-248-30
   US-10-789-273-4
  PRIOR FILING DATE: 2002-01-04
PRIOR APPLICATION NUMBER: 60/259,927
PRIOR FILING DATE: 2001-01-05
NUMBER OF SEQ ID NOS: 60
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 30
   GENERAL INFORMATION:
APPLICANT: Cohen, Bruce D.
APPLICANT: Bebbe, Jean
APPLICANT: Miller, Penelo
APPLICANT: Moyer, James D
   Sequence 50, Application US/11144248 Publication No. US20050244408A1
   Matches
   Query Match
  Query Match
   APPLICANT: COTVAIAN, JOSE R.
APPLICANT: Gallo, Michael
TITLE OF INVENTION: ANTIBODIES TO INSULIN-LIKE GROWTH FACTOR I RECEPTOR
FILE REFERENCE: ABX-PF2
CURRENT APPLICATION NUMBER: US/11/144,248
CURRENT FILING DATE: 2005-06-02
FRIOR APPLICATION NUMBER: US/10/038,591
PRIOR FILING DATE: 2002-01-04
PRIOR FILING DATE: 2002-01-04
PRIOR APPLICATION NUMBER: 60/259,927
PRIOR APPLICATION NUMBER: 60/259,927
PRIOR PILING DATE: 2001-01-05
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   PRIOR APPLICATION NUMBER: US/10/038,591
PRIOR APPLICATION NUMBER: US 10/010,942
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   Miller, Penelope E. Moyer, James D.

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Pred. No. 1.1e-08;
  Score 123; DB 7; Length 473; Pred. No. 4.2e-08;
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문
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  US-11-172-320-6
   US-10-512-184-29
   US-10-512-184-29
  US-10-789-273-4
  APPLICANT: Fraunhofer Gesellschaft zur F"rderung der angewandten Forschung e.V.
TITLE OF INVENTION: Antibodies, recombinant antibodies, recombinant
TITLE OF INVENTION: antibody fragments and fusions mediated plant disease
TITLE OF INVENTION: resistance against fungi
FILE REFERENCE: 3581.01US01
FILE REFERENCE: 3581.01US01
CURRENT APPLICATION NUMBER: US/10/512,184
CURRENT FILING DATE: 2004-10-22
NUMBER OF SEQ ID NOS: 72
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 29
   Sequence 6, Application US/11172320 Publication No. US20050244413A1 GENERAL INFORMATION:
   Sequence 29, Application US/10512184
Publication No. US20050244901A1
GENERAL INFORMATION:
   SEQ ID NO 4
LENGTH: 138
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   Query Match
          APPLICANT: Adolf, Guenther
APPLICANT: Baum, Anke
APPLICANT: Baum, Anke
APPLICANT: Heider, Karl-Heinz
TITLE OF INVENTION: Compositions and Methods for Treating Cancer using
TITLE OF INVENTION: Cytotoxic CD44 Antibody Immunoconjugates and
TITLE OF INVENTION: Chemotherapeutic Agents
FILE REFERENCE: 1/1383
FULL REFERENCE: 1/1383
CURRENT APPLICATION NUMBER: US/11/172,320
CURRENT FILING DATE: 2003-08-21
PRIOR APPLICATION NUMBER: US/10/645,215
PRIOR FILING DATE: 2003-08-21
PRIOR FILING DATE: 2003-08-21
   PRIOR FILING DATE: 2001-12-06
PRIOR APPLICATION NUMBER: US
PRIOR FILING DATE: 2000-12-06
NUMBER OF SEQ ID NOS: 63
  SOFTWARE: FastSEQ for Windows Version 4.0
PRIOR APPLICATION NUMBER: EP 02 018 686.2
  PEATURE:
NAME/KBY: SIGNAL
LOCATION: (1)...(19)
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ORGANISM: Mus musculus
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   77.9%; Score 120; DB 1; Length 263, 76.7%; Pred. No. 5.5e-08; rative 3; Mismatches 4; Indels
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76.7%;
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SEQ ID NO 6

PRIOR FILING DATE: August 21, 2002
PRIOR APPLICATION NUMBER: US 60/405,956
PRIOR FILING DATE: August 26, 2002
NUMBER OF SEQ ID NOS: 9
SOFTWARE: Patentin Ver. 2.1

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; FEATURE:
; OTHER INFORMATION: Humanised Murine Antibody BIWA 4 Heavy Chain
US-11-172-320-6
  US-10-789-273-10
  문
  ; TYPE: PRT; ORGANISM: Homo sapiens US-11-144-248-49
  US-11-144-248-49
  Sequence 49, Application US/11144248 Publication No. US20050244408A1 GENERAL INFORMATION:
  Query Match
Best Local Similarity
Matches 23; Conserv
  SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 49
LENGTH: 470
  Sequence 10, Application US/10789273
Publication No. US20050249725A1
GENERAL INFORMATION:
  Matches
   Query Match
APPLICANT: Basi, Guriq
APPLICANT: Saldanha, Jose
APPLICANT: Saldanha, Jose
APPLICANT: Yednock, Tod
TITLE OF INVENTION: HUMANIZED ANTIBODIES THAT RECOGNIZE
TITLE OF INVENTION: BETA-AMYLOID PEPTIDE
FILE REFERENCE: ELM-002CP
CURRENT APPLICATION NUMBER: US/10/789,273
CURRENT FILING DATE: 2004-02-27
PRIOR APPLICATION NUMBER: US/10/388,389
PRIOR FILING DATE: 2003-03-12
  APPLICANT: Cohen, Bruce D.
APPLICANT: Beebe, Jean
APPLICANT: Miller, Penelor
APPLICANT: Moyer, James D.
  CURRENT APPLICATION NUMBER: US/11/144,248
CURRENT FILLING DATE: 2005-06-02
PRIOR APPLICATION NUMBER: US/10/038,591
PRIOR FILING DATE: 2002-01-04
PRIOR PELICATION NUMBER: 60/259,927
PRIOR APPLICATION NUMBER: 60/259,927
PRIOR FILING DATE: 2001-05
NUMBER OF SEQ ID NOS: 60
  APPLICANT: Corvalan, Jose R.
APPLICANT: Gallo, Michael '
TITLE OF INVENTION: ANTIBODIES TO INSULIN-LIKE GROWTH FACTOR I RECEPTOR
   FILE REFERENCE: ABX-PF2
   TYPE: PRT ORGANISM: Artificial Sequence
   ENGTH: 444
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  Beebe, Jean
Miller, Penelope B.
Moyer, James D.
  QAQLVESGGGLVKPGGSLRLSCAASGFTFS 49
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76.7%;
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76.7%;
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Pred. No. 8.5e-08;
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Pred. No. 1.1e-07;
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   Length 444;
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APPLICANT: Basi, Guriq
APPLICANT: Baldanha, Jose
APPLICANT: Saldanha, Jose
APPLICANT: Saldanha, Tose
APPLICANT: Saldanha, Ted
TITLE OF INVENTION: HUMANIZED ANTIBODIES THAT RECOGNIZE
TITLE OF INVENTION: BETA-AMTICID PEPTIDE
FILE REFERENCE: ELM-002CP
CURRENT APPLICATION NUMBER: US/10/789,273
CURRENT FILING DATE: 2004-02-27
PRIOR APPLICATION UNBERS: US/10/388,389
PRIOR FILING DATE: 2003-03-12
PRIOR APPLICATION NUMBER: US 10/010,942

GENERAL INFORMATION

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RESULT 9
US-11-144-248-32
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US-10-789-273-10
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   Ś
   ; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-144-248-32
                                    US-10-789-273-9
  NUMBER OF SEQ ID NOS: 63
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 10
LENGTH: 98
TYPE: PRT
  GENERAL INFORMATION:
  SEQ ID NO 32
   Sequence 32, Application US/111 Publication No. US20050244408A1
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Sequence 9, Application US/10789273 Publication No. US20050249725A1
   Matches
   Query Match
Best Local Similarity
   Best Local
   APPLICANT: Cohen, Bruce D.
APPLICANT: Beebe, Jean
APPLICANT: Miller, Penelo
  PRIOR APPLICATION NUMBER: US 10/010,942
PRIOR FILING DATE: 2001-12-06
PRIOR APPLICATION NUMBER: US 60/251,892
PRIOR FILING DATE: 2000-12-06
   FILE REFERENCE: ABX-PF2
CURRENT APPLICATION NUMBER: US/11/144,248
CURRENT FILING DATE: 2005-06-02
  PRIOR APPLICATION NUMBER: US/10/038,591
PRIOR FILING DATE: 2002-01-04
PRIOR APPLICATION NUMBER: 60/259,927
PRIOR FILING DATE: 2001-01-05
NUMBER OF SEQ ID NOS: 60
SOFTWARE: Patentin Ver. 2.1
  APPLICANT: Corvalan, Jose R. APPLICANT: Gallo, Michael
  APPLICANT:
  TITLE OF INVENTION: ANTIBODIES TO INSULIN-LIKE GROWTH FACTOR I RECEPTOR
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  Beebe, Jean
Miller, Penelope B.
Moyer, James D.
   Application US/11144248
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  US-10-789-273-9
   US-11-144-248-24
   Sequence 16, Application US/11144248 Publication No. US20050244408A1 GENERAL INFORMATION:
  APPLICANT: Cohen, Bruce D.
APPLICANT: Beebe, Jean
APPLICANT: Miller, Penelope E.
APPLICANT: Moyer, James D.
APPLICANT: Corvalan, Jose R.
APPLICANT: Gallo, Michael
  SOFTWARE: P
SEQ ID NO 24
  Sequence 24, Ap
Publication No.
  Matches
   Query Match
Best Local Similarity
   SEQ ID NO 9
  Matches
  Query Match 76.6%;
Best Local Similarity 73.3%;
   GENERAL INFORMATION:
APPLICANT: Cohen, Bruce D.
APPLICANT: Beebe, Jean
APPLICANT: Miller, Penelope B.
APPLICANT: MILLER, Penelope B.
APPLICANT: Corvalan, Jose R.
APPLICANT: Corvalan, Jose R.
APPLICANT: Gallo, Michael
TITLE OF INVENTION: ANTIBODIES TO INSULIN-LIKE GROWTH FACTOR I RECEPTOR PILE REFERENCE: ABX-PF2
CURRENT APPLICATION NUMBER: US/11/144,248
CURRENT FILING DATE: 2005-06-02
PRIOR APPLICATION NUMBER: US/10/038,591
  CURRENT APPLICATION NUMBER: US/11/144,248
CURRENT FILING DATE: 2005-06-02
PRIOR APPLICATION NUMBER: US/10/038,591
PRIOR FILING DATE: 2002-01-04
   PRIOR FILING DATE: 2001-12-06
PRIOR APPLICATION NUMBER: US 60/251,892
PRIOR FILING DATE: 2000-12-06
NUMBER OF SEO ID NOS: 63
  SOFTWARE: FastSEQ for Windows Version 4.0
  PRIOR FILING DATE: 2001-01-05
NUMBER OF SEQ ID NOS: 60
   PRIOR APPLICATION NUMBER: 60/259,927
  TITLE OF INVENTION: ANTIBODIES TO INSULIN-LIKE GROWTH FACTOR I RECEPTOR FILE REFERENCE: ABX-PF2
  LENGTH: 121
TYPE: PRT
ORGANISM: Homo sapiens
  LENGTH:
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  22;
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   22; Conservative
  122
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  Application US/11144248
No. US20050244408A1
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73.3%;
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Pred. No. 4.8e-08;
   Score 118; DB 1;
Pred. No. 4.8e-08;
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US-10-789-273-12
Sequence 12, Application US/10789273
Publication No. US20050249725A1
GENERAL INFORMATION:
APPLICANT: Basi, Guriq
FAPLICANT: Basi, Guriq
  片
   밁
  US-11-144-248-16
  Ś
   US-10-789-273-8
   RESULT 14
   CURRENT APPLICATION NUMBER: US/10/789,273
CURRENT FILING DATE: 2004-02-27
PRIOR APPLICATION NUMBER: US/10/388,389
PRIOR FILING DATE: 2003-03-12
PRIOR APPLICATION NUMBER: US 10/010,942
PRIOR FILING DATE: 2001-12-06
PRIOR APPLICATION NUMBER: US 60/251,892
PRIOR APPLICATION NUMBER: US 60/251,892
PRIOR PILING DATE: 2000-12-06
  RESULT 13
APPLICANT: Baldanha, Jose
APPLICANT: Saldanha, Jose
APPLICANT: Saldanha, Jose
APPLICANT: Yednock, Ted
TITLE OF INVENTION: HUMANIZED ANTIBODIES THAT RECOGNIZE
TITLE OF INVENTION: BETA-AMYLOID PEPTIDE
FILE REFERENCE: ELN-002CP
CURRENT APPLICATION NUMBER: US/10/789,273
CURRENT APPLICATION NUMBER: US/10/789,273
   Best Loc
Matches
  SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 16
LENGTH: 125
  SEQ ID NO 8
LENGTH: 138
  GENERAL INFORMATION:
   Sequence 8, Application US/10789273
Publication No. US20050249725A1
   Matches
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  Query Match
Best Local
   APPLICANT: Basi, Guriq
APPLICANT: Saldanha, Jose
APPLICANT: Saldanha, Jose
APPLICANT: Yednock, Ted
TITLE OF INVENTION: HUMANIZED ANTIBODIES THAT RECOGNIZE
TITLE OF INVENTION: BETA-AMYLOID PEPTIDE
FILE REFERENCE: ELN-002CP
  PRIOR APPLICATION NUMBER: 60/259,927
PRIOR FILING DATE: 2001-01-05
  NUMBER OF SEQ ID NOS: 63
SOFTWARE: FastSEQ for Windows Version 4.0
  NUMBER OF SEQ ID NOS: 60
   ORGANISM: Homo sapiens
  NAME/KEY: SIGNAL LOCATION: (1)...(19)
  OTHER INFORMATION: Humanized 3D6 heavy chain variable region FEATURE:
   TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
   TYPE: PRT
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   20 EVQLLESGGGLVQPGGSLRLSCAASGFTFS 49
  1 QVQLVQSGGGVVHPGRSLKLSCAGSGFTFS 30 :|||::||||:|||||||
  1 QVQLVQSGGGVVHPGRSLKLSCAGSGFTFS 30 :|||::||||:|||
  Similarity
   Similarity
  EVOLLESGGIVOPGGSLRLSCAASGFTFS 30
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  76.6%;
73.3%;
   76.6%; Score 118; DB 1;
73.3%; Pred. No. 5.3e-08;
bive 5; Mismatches 3
   5
   Score 118; DB 7;
Pred. No. 4.9e-08;
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  Length 125;
   Length 138;
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PRIOR APPLICATION NUMBER: 60/259,927
PRIOR FILING DATE: 2001-01-05
NUMBER OF SEQ ID NOS: 60
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 46
LENGTH: 470
TYPE: PRT
ORGANISM: Homo sapiens
US-11-144-248-46
   ; PRIOR APPLICATION NUMBER: US/10/388,389
; PRIOR FILING DATE: 2003-03-12
; PRIOR APPLICATION NUMBER: US 10/010,942
; PRIOR FILING DATE: 2001-12-06
; PRIOR APPLICATION NUMBER: US 60/251,892
; PRIOR FILING DATE: 2000-12-06
; NUMBER OF SEQ ID NOS: 63
; SOPTWARE: FRAELSEQ for Windows Version 4.0
; SEQ ID NO 12
; LENGTH: 138
  ; FEATURE:
; OTHER INFORMATION: Humanized 3D6 light chain variable region FEATURE:
; NAME/KEY: SIGNAL ; LOCATION: (1)...(19)
US-10-789-273-12
  ₽
Search completed: November 21, 2005, 12:33:52 Job time: 0.649451 secs
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  Sequence 46, Application US/11144248 Publication No. US20050244408A1 GENERAL INFORMATION:
   Query Match 76.6%;
Best Local Similarity 73.3%;
Matches 22; Conservative
   Query Match 76.6%; Score 118; DB 7; Length 470; Best Local Similarity 73.3%; Pred. No. 1.5e-07; Matches 22; Conservative 5; Mismatches 3; Indels
   FILE REFERENCE: ABX-PF2
CURRENT APPLICATION NUMBER: US/11/144,248
CURRENT FILING DATE: 2005-06-02
PRIOR APPLICATION NUMBER: US/10/038,591
PRIOR FILING DATE: 2002-01-04
   APPLICANT: Cohen, Bruce D.
APPLICANT: Beebe, Jean
APPLICANT: Miller, Penelope E.
APPLICANT: Miller, Penelope E.
APPLICANT: Moyer, James D.
APPLICANT: Corvalan, Jose R.
APPLICANT: Gallo, Michael
TITLE OF INVENTION: ANTIBODIES TO INSULIN-LIKE GROWTH FACTOR I RECEPTOR
  TYPE: PRT
ORGANISM: Artificial Sequence
   20
   1 QVQLVQSGGGVVHPGRSLKLSCAGSGFTFS 30
  EVQLLESGGGLVQPGGSLRLSCAASGFTFS 49
   5; Mismatches
  Score 118; DB 1; Length 138; Pred. No. 5.3e-08;
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   0
  0
   Gaps
  Gaps
   0
  0
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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
  Minimum DB
Maximum DB
  Scoring table:
   Sequence:
   Title:
Perfect score:
  OM protein -
   Database :
  Total number of hits satisfying chosen parameters:
   92:
Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
   seq length: 0
seq length: 2000000000
  protein search, using sw model
   BLOSUM62
   US-10-632-706-159
32
  2443163 seqs, 439378781 residues
   Gapop 10.0 , Gapext 0.5
   1 DYDMH 5
   November 21,
   A_Geneseq_21:*
  GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
  geneseqp1980s:*
geneseqp190s:*
geneseqp2000s:*
geneseqp2001s:*
geneseqp2002s:*
geneseqp2003as:*
geneseqp2003as:*
geneseqp2003as:*
   geneseqp2005s:*
   2005, 11:49:32 ; Search time 6.69414 Seconds (without alignments) 328.182 Million cell updates/sec
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## SUMMARIBS

| 113<br>113<br>114<br>115<br>116<br>117<br>117<br>117<br>117<br>117<br>117<br>117<br>117<br>117                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          | Result<br>No.    |
|-----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|------------------|
|                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         | Score            |
| 1000.0                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  | Query<br>Match 1 |
| 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   | Length           |
| <b>ΘΘΘΘωναρουνασυνωυσοριωα</b>                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          | B                |
| ADR38765<br>ADR38767<br>ADR38757<br>AAG00943<br>AAG00943<br>ABG20497<br>ABU70539<br>ABU70539<br>ADS09256<br>ADS44811<br>ABM68679<br>ADN18693<br>ADN18693<br>ADN18693<br>ABB92826<br>AAM93283<br>ABB97214<br>AAY68736<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM93284<br>AAM | ID               |
| Adr38761<br>Adr38761<br>Adr38769<br>Aagr00943<br>Aagr00943<br>Abg20497<br>Abu7053<br>Ads09256<br>Ads44811<br>Abm68679<br>Adn1869<br>Ads41979<br>Adn1869<br>Ads41973<br>Abb9286<br>Ab93283<br>Ab93283<br>Ad1307214<br>Ab9697214<br>Ab9697214<br>Ab9697214<br>Ab9697214<br>Ab9697214<br>Ab969295<br>Adf08477<br>Abm82358463<br>Ads43244<br>Aay69295<br>Ade09984<br>Ade09984<br>Ade09984<br>Ade09984                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       | , p.             |
| Mouse hea Mouse hea Mouse hea Mouse hea Human sec Novel hum Human adi Human c-M Bacterial Bacterial Bacterial Bacterial Human pro Novel hum Short cha Human pro Novel hum Short cha Human MF A human M                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          | on               |

| ADY04516  |
|-----------|
| AAG52810  |
| AAG52811  |
| AAG52812  |
| ABP25940  |
| AAB63286  |
| AAB63457  |
| ABB04486  |
| ABM37298  |
| AAU4 0779 |
| ABP63246  |
| ABJ05221  |
| AAB55829  |
| AAB58101  |
| AAB57691  |
| ADR21541  |
| AAY97696  |
| ABB66508  |
| ABGZU499  |
| ADAS4819  |
|           |
| ABG04416  |

## ALIGNMENTS

ADR38763
ID ADR38763
ID ADR38763
AC ADR33
XX ADR38763
AC ADR3
XX ADR3
AC ADR3
XX MOUS
XX MOUS
XX MOUS
XX MUS20
XX US20
XX US20
XX IS20
antibacterial; antibody; botulinum neurotoxin type A; BoNT/A; BoNT/A neutralising epitope; anti-botulinum neurotoxin antibody; toxin neutralisation; botulinum neurotoxin poisoning; mouse; heavy chain; CDR1; complementarity determinig region 1. 01-AUG-2003; 2003US-00632706. 09-SEP-2004. US2004175385-A1. Mus sp Mouse heavy chain anti-BoNT-antibody CDR1 seqid 167. 02-DEC-2004 ADR38765; ADR38765 standard; peptide; 5 AA. (first entry)

31-AUG-1998; 98US-00144886. 01-AUG-2002; 2002US-0400721P.

New isolated antibody that neutralizes botulinum neurotoxin type A, useful for diagnosing botulism or for treating pathologies associated with botulinum neurotoxin poisoning.

WPI; 2004-652009/63.

Marks JD,

Amersdorfer P;

(REGC ) UNIV CALIFORNIA.

Example 4; SEQ ID NO 167; 110pp; English.

The invention describes an isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a specific clone where (I) binds to and neutralises botulinum neurotoxin type A (BoNT/A). An isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a clone chosen from clone 25, C25, C39, ICG, 3D12, B4, IF3, huC25, A11, Ar2, WR1(Y), WR1(T), 3-1, 3-8, 3-10 and INGI, where (I) binds to and neutralizes botulinum neurotoxin type A (BoNT/A). Also described are: a polypeptide (II) comprising BoNT/A neutralising epitope having an epitope that is

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RESULT 2
ADR33761
ID ADR33
AC ADR3
AC 
  ភិនិសិស្តិសិស្តិសិស្តិសិស្តិសិស្តិសិស្តិសិស្តិ
សិស្តិសិស្តិសិស្តិសិស្តិសិស្តិសិស្តិសិស្តិសិស្តិសិស្តិសិស្តិសិស្តិ
   밁
   ঠ
  specifically bound by an antibody expressed by clones as mentioned in (I) producing (I), and a composition (III) comprising several antibodies, where each antibody is specific for a different epitope of a botulinum neurotoxin and the combination of antibodies shows greater toxin neutralisation than the single antibodies in surplus. The following are disclosed; a pharmaceutical composition comprising (I), and a kit comprising (I). (I) is useful for neutralising BONT/A antibody and for neutralising a botulinum neurotoxin which involves contacting neurotoxin with (I) in surplus, where each of (I) is specific for a different epitope of the botulinum neurotoxin and the combination of antibodies shows greater toxin neutralisation than the single antibodies in surplus. (I) is useful for diagnosing the botulism or for treating pathologies associated with botulinum neurotoxin pathologies associated with botulinum neurotoxin poisoning. (I) exhibits specificity and affinity towards BONT/A. (I) enables rapid detection or diagnosis of botulism. This is the amino acid sequence of mouse heavy chain anti-BONT-antibody CRD1.
   Query Match
Best Local S
Matches 5
  Mue
  antibacterial; antibody; botulinum neurotoxin type A; BONT/A; BONT/A neutralising epitope; anti-botulinum neurotoxin antibody; toxin neutralisation; botulinum neurotoxin poisoning; mouse; heavy chain; CDR1; complementarity determinig region 1.
  New isolated antibody that neutralizes botulinum neurotoxin type A, useful for diagnosing botulism or for treating pathologies associated
   WPI;
  31-AUG-1998; 98US-00144886.
01-AUG-2002; 2002US-0400721P.
  01-AUG-2003; 2003US-00632706.
  09-SEP-2004.
   US2004175385-A1.
  Mouse heavy chain anti-BoNT-antibody CDR1 segid 163
   02-DEC-2004 (first entry)
  ADR38761 standard; peptide; 5 AA.
  Sequence 5 AA;
   (REGC ) UNIV CALIFORNIA.
   2004-652009/63.
  Ą
   l Similarity
5; Conserv
  1 DYDMH 5
  DYDMH 5
  Ameredorfer
   Conservative
  100.0%;
  0; Mismatches
  Score 32;
Pred. No.
  2e+06;
   명
   8,
   Length
  Indels
  0
  Gaps
```

The invention describes an isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a specific clone where (I) binds to and neutralises botulinum neurocxin type A (BoNT/A). An isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a clone chosen from clone specifically bound by an antibody expressed by a clone chosen from clone specifically bound by an antibody expressed by a clone chosen from clone specifically bound by an antibody expressed by a clone chosen from clone specifically bound by an antibody expressed by a clone chosen from clone specifically bound by an antibody expressed by a clone chosen from clone specifically with the specifically binds to and neutralizes botulinum

The invention describes an isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a specific clone where (I) binds to and neutralises botulinum neurotoxin type A (BoNT/A). An isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a clone chosen from clone

Example 4;

SEQ ID NO 163; 110pp; English.

neurotoxin

poisoning.

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ADRIBULT 3
ADRIBUT 3
ADRIBUT 1
ADRIBUT ADRIBUT
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  ទន្ទន្ទន្ទន្ទន្ទន្ទន្ទន្ទន្ទន្ទន្ទន្ទន្
  문
   CC comprising BONT/A neutralising epitope having an epitope that is comprising BONT/A neutralising epitope having an epitope that is comprising Bont/A neutralising epitope having an epitope that is comprising several anti-
CC producing (I); and a composition (III) comprising several anti-
CC different epitope of a botulinum neurotoxin and the combination of continuous products and the combination of antibodies howe greater toxin neutralisation than the single antibodies in surplus. The following are disclosed: a pharmaceutical composition comprising (I); and a kit comprising (I). (I) is useful for neutralising CC emprising (I) and a kit comprising (I). (I) is useful for neutralising composition of comprising (I) and a kit comprising (I) and useful for neutralising composition of comprising (I) antibody and for neutralising a botulinum neutrotoxin which comprising (I) is useful for neutralising composition of (I) is single antibodies in surplus. (I) is surplus, where each of (I) is single antibodies in surplus. (I) is useful for diagnosing the botulism corrections of antibodies associated with botulinum neutrotoxin containts or for treating pathologies associated with botulinum neutrotoxin containts poisoning. (I) exhibits specificity and affinity towards BoNT/A. (I) cenables rapid detection or diagnosis of botulism. This is the amino acid or sequence of mouse heavy chain anti-BoNT-antibody CRDI.
   Query Match
Best Local S
Matches 5
   WPI;
   antibacterial; antibody; botulinum neurotoxin type A; BONT/A; BONT/A neutralising epitope; anti-botulinum neurotoxin antibody; toxin neutralisation; botulinum neurotoxin poisoning; mouse; heavy chain; CDR1; complementarity determinig region 1.
   New isolated antibody that neutralizes botulinum neurotoxin type A, useful for diagnosing botulism or for treating pathologies associated
  31-AUG-1998; 98US-00144886.
01-AUG-2002; 2002US-0400721P.
  Sequence 5 AA;
   Marks
  01-AUG-2003; 2003US-00632706
  09-SEP-2004
  US2004175385-A1
   gs suM
   02-DEC-2004
   ADR38757;
   ADR38757 standard;
  Example
   (REGC ) UNIV
   Mouse
  botulinum
   2004-652009/63.
   à
  heavy
  1 DYDMH 5
   5; Conserv
   4
   SEQ
   Ameredorfer
  chain anti-BoNT-antibody CDR1 seqid 159.
   100.0%;
ilarity 100.0%;
Conservative 0,
   (first entry)
  CALIFORNIA
   ij
   neurotoxin poisoning.
   ð
   peptide; 5
  159; 110pp; English
   P;
  ٥,
   Score 32; DB
Pred. No. 2e+
0; Mismatches
   2e+06;
  DB 8;
   0
   Length 5;
   0
   Gaps
   0,
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  CC 3-8, 3-10 and ING1, where (I) binds to and neutralizes botulinum CC neurotoxin type A (BoNT/A). Also described are: a polypeptide (II) CC comprising BoNT/A neutralising epitope having an epitope that is CC specifically bound by an antibody expressed by clones as mentioned in (I) Cf. producing (I); and a composition (III) comprising several anticodies, where each antibody is specific for a different epitope of a botulinum neurotoxin antibodies, where each antibodies described are single antibodies in surplus. The following are disclosed; a pharmaceutical composition (CC intibodies shows greater toxin neutralisation than the single antibodies (I); and a kit comprising (I) is useful for neutralising comprising (I); and a kit comprising (I). (I) is useful for neutralising comprising (I) and a composition with (I) in surplus, where each of (I) is specific for a different epitope of the botulinum neurotoxin and the combination of antibodies shows greater toxin neutralisation than the CC single antibodies in surplus. (I) is useful for diagnosing the botulism CC or for treating pathologies associated with botulinum neurotoxin conduction or diagnosis of botulism. This is the amino acid composes rapid detection or diagnosis of botulism. This is the amino acid compose of mouse heavy chain anti-BoNT-antibody CRD1.
  Matches
   Query Match
   Sequence 5 AA;
   antibacterial; antibody; botulinum neurotoxin type A; BONT/A; BONT/A neutralising epitope; anti-botulinum neurotoxin antibody; toxin neutralisation; botulinum neurotoxin poisoning; mouse; heavy chain; CDR1; complementarity determinig region 1.
   Mouse heavy chain anti-BoNT-antibody CDR1 seqid 171.
  ADR38769 standard; peptide; 5 AA.
   New isolated antibody that neutralizes botulinum neurotoxin type A, useful for diagnosing botulism or for treating pathologies associated
   Marks JD,
   31-AUG-1998; 98US-00144886.
01-AUG-2002; 2002US-0400721P.
  01-AUG-2003; 2003US-00632706.
  US2004175385-A1.
   02-DEC-2004 (first entry)
  WPI; 2004-652009/63.
  (REGC ) UNIV CALIFORNIA.
   Local Similarity 100
les 5; Conservative
  botulinum neurotoxin poisoning.
   1 DYDMH 5
   Amersdorfer P;
  100.0%;
  0; Mismatches
  Score 32; DB 8;
Pred. No. 2e+06;
  DB 8;
  0
  Length 5;
  Indels
  0
  Gaps
  0
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The invention describes an isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a specific clone where (I) binds to and neutralises botulinum neurotoxin type A

The present sequence is a polypeptide encoded by one of a large number of 5' ESTs derived from mRNAs encoding secreted proteins. The 5' ESTs were prepared from total human RNAs or polyA+ RNAs derived from 30 different

Example 4;

SEQ ID NO 171; 110pp; English

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RESULT 5
AAG00943
AID 0943
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  CC (BoNT/A). An isolated antibody (I) that specifically binds to an epitope CC specifically bound by an antibody expressed by a clone chosen from clone CS 25, C25, C39, 1C6, 3D12, B4, 1F3, huC25, Ar1, KR1(V), WR1(T), 3-1, CC comprising BONT/A neutralising epitope dare: a polypeptide (II) CC comprising BONT/A neutralising epitope having an epitope that is CC specifically bound by an antibody expressed by clones as mentioned in (I) producing (I); and a composition (III) comprising several anticce different epitope of a botulinum neurotoxin antibodies, where each antibody is specific for a CC different epitope of a botulinum neurotoxin and the combination of CC comprising (I); and a kit comprising (I) is useful for neutralising a botulinum neurotoxin which CC comprising (I); and a kit comprising (I) is useful for neutralising ENOT/A antibody and for neutralising a botulinum neurotoxin which CC involves contacting neurotoxin with (I) in surplus, where each of (I) is specific for a different epitope of the botulinum neurotoxin and the combination of antibodies shows greater toxin neutralisation than the single antibodies in surplus. (I) is useful for diagnosing the botulism or for treating pathologies associated with botulinum neurotoxin cc enables rapid detection or diagnosis of botulism. This is the amino acid companies of mouse heavy chain anti-BoNT-antibody CRD1.
  밁
   S
   Matches
  Query Match
   WPI; 200
N-PSDB;
  Claim 13; SEQ ID NO 5024; 71pp + Sequence Listing; English.
   New nucleic acid that is a 5' expressed sequence tag (5' EST) for obtaining cDNAs and genomic DNAs that correspond to 5'ESTs and fo diagnostic, forensic, gene therapy and chromosome mapping procedu
   EP1033401-A2
   Human; 5' EST; expressed sequence tag; secreted protein; cDNA isolation; gene therapy; chromosome mapping.
   Human secreted
   06-OCT-2000
  Sequence 5 AA;
   Dumas Milne Edwards J,
  26-FEB-1999;
  21-FEB-2000; 2000EP-00200610.
  06-SEP-2000.
   Homo sapiens.
   AAG00943;
   AAG00943 standard; protein; 142
   (GEST ) GENSET
   Local Similarity
   2000-500381/45.
DB; AAC00949.
   DYDMH 5
   100.0%; Score 32; DB 8; ilarity 100.0%; Pred. No. 2e+06; Conservative 0; Mismatches 0
   (first entry)
   protein, SEQ ID NO: 5024.
   99US-0122487₽
  Duclert A,
   Giordano
   0
   Length 5;
   mapping procedures
   Indels
   0
   Gaps
  0
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RESULT 6
ABG20497
ID ABG2
XX ABG2
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XX ABG2
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  tissues. EST sequences usually correspond mainly to the 3' untranslated region (UTR) of the mRNA because they are often obtained from oligo-dT primed cDNA libraries. Such ESTs are not well suited for isolating cDNA sequences derived from the 5' ends of mRNAs and even in those cases where longer cDNA sequences have been obtained, the full 5' UTR is rarely included. 5' ESTs are derived from mRNAs with intact 5' ends and can therefore be used to obtain full length cDNAs and genomic DNAs. 5' ESTs are also used in diagnostic, forensic, gene therapy and chromosome mapping procedures. They are used to obtain upstream regulatory sequences and to design expression and secretion vectors
  New isolated polynucleotide and encoded polypeptides, useful in diagnostics, forensics, gene mapping, identification of mutatio responsible for genetic disorders or other traits and to assess
   Claim 20; SEQ ID NO 50856; 103pp; English.
  N-PSDB; AAS84684.
  Drmanac
  31-MAR-2000; 2000US-00540217.
23-AUG-2000; 2000US-00649167.
   30-MAR-2001; 2001WO-US008631
   WO200175067-A2
  Human; chromosome mapping; gene mapping; gene therapy; forensic; food supplement; medical imaging; diagnostic; genetic disorder.
  Novel human diagnostic protein #20488
   18-FEB-2002
   ABG20497;
  ABG20497 standard; protein; 144 AA
  Sequence 142 AA;
   (HYSE-) HYSEQ
   sapiens
  2001-639362/73
  22
  1 DYDMH 5
  Similarity
5; Conserv
  RT,
  DYDMH 26
   Conservative
   Liu C,
  (first entry
   INC.
  100.0%;
   ΤŻ
  0
  Score 32;
Pred. No.
  Mismatches
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   0
   Length 142
  Indels
  mutations
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  Gaps
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sequences. (I) is useful as hybridisation probes, polymerase Chain reaction (PCR) primers, oligomers, and for chromosome and gene mapping, and in recombinant production of (II). The polymucleotides are also used in diagnostics as expressed sequence tags for identifying expressed genes. (I) is useful in gene therapy techniques to restore normal activity of (II) or to treat disease states involving (II). (II) is useful for generating antibodies against it, detecting or quantitating a polypeptide in tissue, as molecular weight markers and as a food supplement. (II) and its binding partners are useful in medical imaging of sites expressing (II). (I) and (II) are useful for treating disorders involving aberrant protein expression or biological activity. The polypeptide and polynucleotide sequences have applications in

protein

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comprising

the polypeptides

and

The invention relates to isolated polynucleotide (I) and polypeptide (II)

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                   adipocyte cells, given in the specification. The proteins are identified by selecting a bait protein from a known adipocyte marker and then performing a yeast 2-hybrid selection to isolate prey proteins encoded by members of an adipocyte cDNA library. The proteins are designated SID (RTM) (selected interacting domains) proteins. Also included are a polymuclectide encoding a polypeptide in the adipocyte cells, a recombinant host cell expressing at least one of the interacting polypeptides of the complex, selecting a modulating compound in adipocyte cells, a SID (RTM)) polypeptide comprising any of the 738 amino acid sequences given in the specification (including its fragment or variant), a vector comprising the SID (RTM) polynuclectide comprising any of the 738 nucleotide sequences given in the specification (including its fragment or variant), a vector comprising the SID (RTM) polynuclectide, a recombinant host cell
  diagnostics, forensics, gene mapping, identification of mutations responsible for genetic disorders or other traits to assess biodiversity and to produce other types of data and products dependent on DNA and amino acid sequences. ABG00010-ABG30377 represent novel human diagnostic mino acid sequences of the invention. Note: The sequence data for this patent did not appear in the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences
  Human; prey; adipocyte; SID; selected interacting domain; antidiabetic; protein-protein interaction; diabetes; yeast 2-hybrid assay; metabolic disorder; obesity.
  New complex between two interacting proteins in adipocyte cells, for identifying selected interacting domains that modulate proteinteractions, or for preventing or treating metabolic disorders a
  Human
  Sequence 144 AA;
   The invention relates to a complex between two interacting
  Claim
  N-PSDB;
   14-MAR-2001; 2001US-0275734P
  14-MAR-2002; 2002WO-EP003768.
   WO200286122-A2.
   10-JUN-2003
  ABU70539 standard;
   (HYBR-) HYBRIGENICS.
  Homo sapiens.
   2003-103412/09.
DB; ACA57083.
  6; Page 163; 382pp; English.
   adipocyte Selected Interacting
   26 DYDMH 30
  1 DYDMH 5
   ۵,
  5
   or diabetes.
   Similarity
  Daviet L;
  Conservative
   (first entry)
  protein; 180
   100.0%;
  0
  Score 32;
Pred. No.
   Mismatches
   domain,
   64;
   BB
   4.
  SID,
  0
   Length 144;
   anorectic;
  proteins
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   record comprising all or part of the data, listed in the specification. The complex, polypeptides, polynucleotides and compounds are useful for preventing or treating metabolic disorders such as obesity or diabetes. The polynucleotides are useful as probes or primers. The complex is particularly useful for identifying selected interacting domains (SID (RTM)) for screening drugs that modulate the protein interaction, thus exhibiting the therapeutic effect. The present sequence represents a SID (prey) protein of the invention
The invention relates to a novel c-Met protein tyrosine kinase antibody or its antigen binding portion that specifically binds to c-Met. The c-Met antibody comprises any one of 1-60 fully defined sequence of 238, 244, 240, 250, 251, 242, 245, 247, 246, 253, 249, 243, 241, etc., amino acids as given in the specification, or its fragment. The invention further comprises: a pharmaceutical composition comprising the c-Met protein tyrosine kinase antibody and a carrier; an isolated cell that produces the c-Met protein tyrosine kinase antibody; and an isolated mucleic acid molecule that comprises a nucleic acid sequence that encodes a heavy chain or its antigen-binding portion of the c-Met protein tyrosine kinase antibody. The c-Met protein tyrosine kinase antibody has cytostatic,
  c-Met; tyrosine kinase antibody; antigen binding; cytostatic; ophthalmological; antiinflammatory; analgesic; vasotropic; antipsoriatic; osteopathic; cancer; tumour; ophthalmic disease; glaucoma; retinitis; retinopathy; uvesitis; ocular photophobia; macular degeneration; pain; acute injury; eye; hyperproliferative disorder; restenosis; angioplasty;
   Novel c-Met protein tyrosine kinase antibody or its antigen-binding portion specifically binding to c-Met, useful for manufacture of medicament for treating cancer or tumor and for treatment of ophthalmic diseases such as glaucoma.
  Human c-Met protein tyrosine kinase antibody, PGIA-2-A4
   Sequence 180 AA;
  Claim 1; SEQ ID NO 18; 303pp; English.
  N-PSDB; ADS09316.
   WPI; 2004-616044/59.
   Morton
   13-FEB-2003; 2003US-0447073P.
   11-FEB-2004;
   WO2004072117-A2
   Homo sapiens.
  psoriasis; HGF; osteoporosis; cancer.
   18-NOV-2004 (first entry)
  ADS09256;
  ADS09256 standard;
  (PHAA ) PHARMACIA CORP
  22
   PA,
   1 DYDMH 5
  ş
  Similarity
  DYDMH 26
  Conservative
   Arbuckle JA,
   2004WO-IB000503.
  protein;
  100.0%;
   Evans ML,
  0
   Score 32; DB
Pred. No. 81;
0; Mismatches
   Joy WD,
  6; Length 180;
   Kahn
  0
   ĽB,
   Shieh JJ
  0
  Gaps
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  cc ophthalmological, antiinflammatory, analgesic, vasotropic, antipsoriatic, cand osteopathic activities. The c-Met protein tyrosine kinase antibody is ct tumour. The c-Met protein tyrosine kinase antibody is ct tumour. The c-Met protein tyrosine kinase antibody is useful for concer or cc diagnosing the presence or ligation of c-Met expressing tissue. The c-Met protein tyrosine kinase antibody is useful for detecting c-Met in a cc protein tyrosine kinase antibody is useful for detecting c-Met in a cc biological sample in vitro or in vivo. The c-Met protein tyrosine kinase cc diseases such as glaucoma, retinitis, retinopathies (e.g., diabetic cretinopathy), uveitis, ocular photophobia, macular degeneration and pain casociated with acute injury to the eye. The pharmaceutical composition is useful for the treatment of hyperproliferative disorders such as crestenosis after angioplasty, and psoriasis, and for the treatment of canimals that lack sufficient HGF, e.g. osteoporosis and cancer. This sequence represents the protein of a phage display generated human c-Met cc antibody of the invention.
  Matches
   Query Match
Best Local
   Recombinant DNA construct; transformed plant; improved plant property; cold tolerance; heat tolerance; drought tolerance; herbicide; osmosis; pathogen tolerance; pest tolerance; push tolerance; pathogen tolerance; pest tolerance; plant disease resistance; cell cycle pathway modification; plant growth regulator; homologous recombination; seed oil yield; protein yield; carbohydrate; homologous recombination; seed oil yield; protein yield; carbohydrate;
  Cao
   Sequence 253 AA;
   21-FEB-2002; 2002US-0360039P
   20-FEB-2003; 2003US-00369493.
   18-DEC-2003
   Bacteria
  Bacterial polypeptide #23241.
  02-DEC-2004
  ADS44811;
   ADS44811 standard; protein; 319
  (CAOY/)
  US2003233675-A1
  bacterial
  (SLAT/)
  31 DYDMH 35
  1 DYDMH 5
  ) CAO Y.
) HINKLE G J.
) SLATER S C.
) CHEN X.
) GOLDMAN B S.
   Similarity
5; Conserv
  Hinkle GJ,
  phosphorus; photosynthesis; lignin; galactomannan;
polypeptide.
  Conservative
  (first entry)
   Slater SC,
   100.0%;
  0
   Score 32; DB 8;
Pred. No. 1.2e+02;
  Mismatches
   Chen
   ×
   Goldman
  Length 253;
   Indels
   Gaps
   0
```

New recombinant DNA construct comprising a promoter positioned to provide for expression of a polynucleotide encoding a polypeptide from a microbial source, useful for producing plants with improved properties.

WPI; 2004-061375/06

The invention relates to a recombinant DNA construct comprising

Claim 1;

SEQ

ID NO 23241; 122pp; English

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ABMG 8679

ID ABMG
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AC ABMG
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  CC provide for expression of a polynucleotide encoding a polypeptide from a CC microbial source. The invention also relates to a transformed plant CC comprising the recombinant DNA construct and a method of producing a CC transformed plant having an improved property. The plant is a crop plant CC such as maize or soybean. The method of producing a transformed plant CC having an improved property comprises transformed plant is a crop plant CC having an improved property comprises transformed plant, where the CC polynucleotide or polypeptide is useful for improving plant properties. CC improved plant properties, e.g. improved cold, heat or drought tolerance, increased resistance to plant disease, better growth rate by modification CC increased resistance to plant disease, better growth rate by modification of the cell cycle pathway with plant growth regulators, increased rate of CC content, improved yield by modification of protein yield and/or CC content, improved plant growth and development under at least one stress CC condition, improved lignin production of photosynthesis or by CC providing improved plant growth and development under at least one stress CC condition, improved lignin production of improved galactomannan CC production. Note: The sequence data for this patent did not CC form part of the printed specification but was obtained in electronic CC format from USPTO at seqdata.uspto.gov/sequence.html.
  Matches
   Query Match
Best Local &
  Antibacterial; fungicide; insecticide; polymorphism; genetic analysis; detection; food; gene expression; plant; animal; microorganism; toxin; antibiotic; biopesticide; virulence factor; disease model; plague;
   Claim
  useful e.g.
  Genomic sequence of Photorhabdus luminescens and encoded polypeptides, useful e.g. as therapeutic antimicrobials and agricultural pesticides.
  WPI; 2003-148459/14.
  Duchaud E,
Buchrieser
  07-FEB-2001; 2001FR-00001659
  07-FEB-2002; 2002WO-IB003040.
  WO200294867-A2
  Photorhabdus luminescens
   whooping cough.
   Photorhabdus luminescens protein sequence #1776.
  20-NOV-2003 (first entry)
  ABM68679;
   ABM68679 standard; protein; 341 AA
   Sequence 319 AA;
   195 DYDMH 199
   2
  l Similarity
5; Conserve
  1 DYDMH 5
  INST PASTEUR.
CNRS CENT NAT RECH SCI.
   SEQ ID
  Conservative
  Taourit S,
   NO 1776; 1205pp; French
   100.0%;
  Glaser P,
  0; Mismatches
   Score 32;
Pred. No.
  Frangeul L,
  1.5e+02;
   DB 8;
   Length 319;
   Kunst
  Indels
   Danchin
  0,
  Gaps
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RESULT 11
ADS41979
ID ADS41979
ID ADS411979
IT ADS411979
IT O2-DE
XX ADS41
XX ADS41
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   CC The invention relates to the isolation of genes and their encoded CC proteins from Photorhabdus luminescens. The isolated sequences are CC sources of probes and primers for detecting the genome of P. luminescens CC and related species, to study polymorphisms, for gene analysis and for CC detection/amplification of the genes. Antibodies (Ab) raised against the CC polypeptides encoded by the genes are used for detection/identification CC of P. luminescens, e.g. in foods. The genes, proteins, Ab and cells that CC carry a gene-containing vector are used to select compounds that CC modulate, induce or inhibit expression of the genes in plants, CC animals or microorganisms other than P. luminescens and are able to alter response or sensitivity to toxins and antibiotics produced by P. CC luminescens. Cells transformed to express the genes are useful for CC entibacterials useful as insecticides, bacterides and fungicides. The genes, proteins, vectors containing the genes and Ab are also useful CC antibacterially (to treat microbial infection by bacteria or fungi that the sepseutically (to treat microbial infection by bacteria or fungi that CC are sensitive to P. luminescens-encoded toxins or antibiotics) and as biopositicides. Other uses of the genes and the proteins are as virulence CC factors and for identifying targets of human diseases for which P. CC luminescens is a model (particularly plague and whooping cough). This sequence represents one of the isolated P. luminescens proteins
  Query Match
Best Local Similarity
  Matches
  Recombinant DNA construct; transformed plant; improved plant property; cold tolerance; heat tolerance; drought tolerance; herbicide; osmosis; pathogen tolerance; pest tolerance; plant disease resistance; cell cycle pathway modification; plant growth regulator; homologous recombination; seed oil yield; protein yield; carbohydrate; nitrogen; phosphorus; photosynthesis; lignin; galactomannan; bacterial polypeptide.
  Sequence 341 AA;
  02-DEC-2004 (first entry)
  ADS41979 standard; protein; 457
   Cao Y,
   (HINK/)
  21-FEB-2002; 2002US-0360039P
   20-FEB-2003; 2003US-00369493.
  US2003233675-A1
  Bacteria.
   Bacterial polypeptide #20409
   WPI; 2004-061375/06.
   18-DEC-2003
   (GOLD/)
  (CAOY/)
  194 DYDMH 198
  HINKLE G J.
) SLATER S C.
) CHEN X.
) GOLDMAN B S.
  1 DYDMH 5
  <u>ت</u>
   Hinkle GJ,
  CAO Y.
  Conservative
   Slater
  100.0%;
   SC,
  0
  Score 32; DB
Pred. No. 1.6
0; Mismatches
   Chen
   ×
   멂
   Goldman
  .6e+02;
   6
   Length 341;
   Indels
   <u>.</u>
   Gaps
   0
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New recombinant DNA construct comprising a promoter positioned to provide for expression of a polynucleotide encoding a polypeptide from a

Claim 1; SEQ ID NO 20409; 122pp; English.

microbial source, useful for producing plants with improved properties.

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   Recombinant DNA construct; transformed plant; improved plant property; cold tolerance; heat tolerance; drought tolerance; herbicide; osmosis; pathogen tolerance; pest tolerance; plant disease resistence; cell cycle pathway modification; plant growth regulator; homologous recombination; seed oil yield; protein yield; carbohydrate; nitrogen; phosphorus; photosynthesis; lignin; galactomannan; bacterial polypeptide.
 (CAOY/)
(HINK/)
(SLAT/)
(CHEN/)
(GOLD/)
  Bacterial polypeptide #1346.
   02-DEC-2004
   ADN18693;
  ADN18693 standard; protein; 465 AA.
  21-FEB-2002; 2002US-0360039P
  20-FEB-2003; 2003US-00369493.
   18-DEC-2003
   US2003233675-A1
   235 DYDMH 239
CAO Y.
HINKLE G J.
SLATER S C.
CHEN X.
GOLDMAN B S.
  (first entry)
```

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The invention relates to a recombinant DNA construct comprising a CC promoter functional in a plant cell, where the promoter is positioned to CC provide for expression of a polynucleotide encoding a polypeptide from a CC microbial source. The invention also relates to a transformed plant CC transformed plant baving an improved property. The plant is a crop plant CC such as maize or soybean. The method of producing a transformed plant CC having an improved property comprises transforming a plant with the recombinant DNA construct and growing the transformed plant, where the CC polynucleotide or polypeptide is useful for improving plant with the recombinant DNA construct is useful for improving plant properties. The recombinant properties, e.g. improved cold, heat or drought tolerance, tolerance to herbicides; extreme osmotic conditions, pathogens or pests, increased resistance to plant disease, better growth rate by modification CC improved plant properties with plant growth regulators, increased rate of content, improved yield by modification of carbohydrate, nitrogen or phosphorus use and/or uptake, by modification of potcein yield and/or CC condition, improved lignin production or improved galactomannan CC condition, This sequence represents a bacterial polypeptide used in the growt of the printed specification but was obtained in electronic CC format from USPTO at sequence data for this patent did not correctly constructed and sequence.
   RESULT 13
ABB92826
  The invention relates to a recombinant DNA construct comprising a CC promoter functional in a plant cell, where the promoter is positioned to CC provide for expression of a polynuclectide encoding a polypeptide from a CC microbial source. The invention also relates to a transformed plant comprising the recombinant DNA construct and a method of producing a transformed plant comparising an improved property. The plant is a crop plant construct and growing an transformed plant construct and growing a transformed plant construct and growing the transformed plant construct and growing the transformed plant where the polynuclectide or polypeptide is useful for improving plant properties.

CC The recombinant DNA construct is useful for producing plant properties. The recombinant DNA construct is useful for producing plant properties. The recombinant DNA construct is useful for producing plant properties. The recombinant DNA construct is useful for producing plants with construct in the recombination of pathways plant proved cold, heat or drought tolerance, content, improved plant disease, better growth rate by modification of the cell cycle pathway with plant growth regulators, increased rate of content, improved yield by modification of carbohydrate, nitrogen or content, improved plant growth and development under at least one stress condition, improved plant growth and development under at least one stress condition, improved lignin production or improved galactomannan cc production. Note: The sequence data for this patent did not form part of the printed specification but was obtained in electronic format from USPTO at sequence.html.
밁
  Ś
   Query Match
Best Local &
   Matches
   New recombinant DNA construct comprising a promoter positioned to provide for expression of a polynucleotide encoding a polypeptide from a microbial source, useful for producing plants with improved properties.
  Cao
   Herbicidal; plant; agriculture; herbicide
   Herbicidally active polypeptide SEQ ID NO 2037.
  Claim 1; SEQ ID NO 1346; 122pp; English.
   WPI; 2004-061375/06.
                         Tietjen K, Weidler
   28-AUG-2001; 2001WO-EP009892
  28-AUG-2001; 2001WO-EP009892.
   07-FEB-2002.
   WO200210210-A2
   Arabidopsis thaliana.
   31-MAY-2002
  ABB92826;
   ABB92826 standard; protein; 473 AA.
   Sequence 465 AA;
   (FARB ) BAYER AG
  Local Similarity
   241 DYDMH 245
  1 DYDMH 5
  Hinkle GJ, Slater
   5
  Conservative
   (first entry
                            3
  100.0%;
  SC,
   0,
  Score 32; DB 8; Pred. No. 2.2e+02;
   Chen
  Mismatches
  ×
   Goldman
  0
   Length 465;
  Indels
  0
  Gaps
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Matches Query Match Best Local

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Similarity

100.0%; ilarity 100.0%; Conservative 0

0

Score 32; DB 8; Length 457; Pred. No. 2.2e+02; O; Mismatches 0; Indels

0;

Gaps

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1 DYDMH 5

Sequence 457 AA;

2002-269010/31

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RESULT 14
AAM93283
ID AAM93
XX AAM93
XX AAM93
XX AAM93
XX AC Human
XX Human
XX Human
XX Homo
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Matches 5
   for herbicidally active compounds, comprising aligning and comparing nucleic acid or amino acid sequences from plant with nucleic acid or amino acid sequences from non-plant organisms using suitable search parameters, where plant sequences having an E-value greater by a factor of 3 than the E-value of most similar non-plant sequences are selected. The polypeptides or nucleic acids encoding them are useful for identifying modulators. The identified modulators are useful as
     The invention relates to primers for synthesising full length cDNA clones. 830 cDNA molecules encoding a human protein have been isolated and nucleotide sequences of 5'- and 3'-ends of the cDNA molecules have been determined. Primers for synthesising the full length cDNA are useful for clarifying the function of the protein encoded by the cDNA. The full length clones were obtained by construction of full length enriched cDNA
  organisms.
   Identifying plant target proteins for herbicidally active compounds, comprising aligning and comparing nucleic acid or amino acid sequences from plant with nucleic acid or amino acid sequences from non-plant
   08-JUL-1999; 99JP-00194486.
11-JAN-2000; 2000JP-00118774.
02-MAY-2000; 2000JP-00183765.
   Claim
  Disclosure;
   830 Primers useful for synthesizing in genetic manipulation.
   Wakamatsu
   07-JUL-2000; 2000EP-00114089
  EP1130094-A2
   Homo sapiens
  Human polypeptide,
   06-NOV-2001
   AAM93283
  AAM93283
  Sequence 473 AA;
  The invention relates to identifying target proteins (ABB90790-ABB94016)
   62
   5; SEQ ID
   full length
   1 DYDMH 5
   Similarity
5; Conserv
  Nishikawa T, :
su A, Sugiyama
   HELIX RES INST
  standard;
   DYDMH 66
  SEQ ID NO
  Conservative
   (first
  ö
   CDNA;
  SEQ ID NO:
   protein;
   2037;
   entry
   100.0%;
  Isogai T,
na T, Nagai
  2764; 1380pp +
  cDNA synthesis; oligo-capping
   261pp +
  493
  0,
   Score 32;
Pred. No.
  2764.
   Mismatches
  Hayashi K,
K, Kojima
   Sequence Listing; English
   full length
  Sequence Listing;
   DB 5; 1
   Ishii S,
S, Otsuki
  Length 473
  CDNA
   clones
   Kawai Y;
T, Koga
  English
   and
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   Ħ;
   Gaps
   0
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Query Match Best Local S Matches 5

Similarity 5; Conserv

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Score 32; DB Pred. No. 2.4 0; Mismatches

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Length 493;

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RESULT 1
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Matches 5
   08-JUL-1999;
11-JAN-2000;
02-MAY-2000;
07-JUL-2000;
  libraries that were synthesised by the oligo-capping method. The primers enable the production of the full length cDNA easily without any special methods. The present sequence is a polypeptide provided in the specification. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in CD-ROM format directly from EPO
   This invention relates to a novel primer's useful for synthesising full length cDNA molecules that encode human proteins. Specifically, it refers to secretory or membrane proteins that are potential therapeutic agents/ target molecules in the field of medicine, and in particular genes encoding proteins that are associated with signal transduction, glycoproteins and transcription. The present invention describes a method for efficiently cloning a full length human cDNA from both the 5° and 3° ends using the oligo-capping method. This polypeptide sequence is a full length human protein of the invention.
   New oligonucleotide primers length human cDNAs.
   07-JUL-2000; 2003EP-00025638
   EP1396543-A2
   oligo-capping
   Human
  Sequence 493
  Example 1; SEQ ID NO 2764; 1340pp; English.
   10-MAR-2004
  human; medicine; signal transduction; glycoprotein; transcription;
  (REAS-)
   ADL30731 standard;
   15
   2004-204755/20.
)B; ADL30730.
  protein
  50 DYDMH
   1 DYDMH
   RES ASSOC BIOTECHNOLOGY
   Nishikawa T,
  Similarity
   2000JP-00118774.
2000JP-00183865.
2000EP-00114089.
  A
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  (first
   encoded
  Sugiyama
   σı
  99JP-00194486
  protein;
   100.0%;
  by a
   Isogai T,
   Nagai K,
  full length cDNA clone SeqID 2764.
  493
  0
  (830 cDNAs) useful
  Score 32; DB
Pred. No. 2.4
); Mismatches
   Hayashi K,
K, Kojima
  2.4e+02;
   DB
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   4
   Ishii S,
   for
   Length 493;
   synthesizing
  Kawai
T, Ko
  0
  Gaps
   Ξ
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| Q                                                                | GenCore version 5.1.6<br>Copyright (c) 1993 - 2005 Compugen Ltd.                                                      |
|------------------------------------------------------------------|-----------------------------------------------------------------------------------------------------------------------|
| OM protein - protei                                              | OM protein - protein search, using sw model                                                                           |
| Run on: Nov                                                      | November 21, 2005, 11:49:36 ; Search time 1.21795 Seconds<br>(without alignments)<br>394.995 Million cell updates/sec |
| Title: US-<br>Perfect score: 32<br>Sequence: 1 D                 | US-10-632-706-159<br>32<br>1 DYDMH 5                                                                                  |
| Scoring table: BLO Gap                                           | BLOSUM62<br>Gapop 10.0 , Gapext 0.5                                                                                   |
| Searched: 283                                                    | 283416 segs, 96216763 residues                                                                                        |
| Total number of hit                                              | Total number of hits satisfying chosen parameters: 283416                                                             |
| Minimum DB seq length: 0 Maximum DB seq length: 200000000        | th: 0<br>th: 2000000000                                                                                               |
| Post-processing: Minimum Match<br>Maximum Match<br>Listing first | Minimum Match 0%<br>Maximum Match 100%<br>Listing first 45 summaries                                                  |
| Database : PI<br>1:<br>2:<br>3:                                  | PIR 80:* : pir1:* : pir2:* : pir3:* : pir4:*                                                                          |

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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| 29         | 28        | 27           | 26           | 25     | 24           | 23         | 22           | 21           | 20           | 19                | 18               | 17           | 16         | 15         | 14     | 13           | 12                 | 11         | 10          | 9           | 80           | 7            | 6           | υ            | 4.         | ω        | N          |            | Result<br>No.  |
|------------|-----------|--------------|--------------|--------|--------------|------------|--------------|--------------|--------------|-------------------|------------------|--------------|------------|------------|--------|--------------|--------------------|------------|-------------|-------------|--------------|--------------|-------------|--------------|------------|----------|------------|------------|----------------|
| 28         | 28        | 28           | 28           | 28     | 28           | 28         | 28           | 28           | 28           | 28                | 28               | 29           | 29         | 29         | 29     | 29           | 29                 | 29         | 29          | 29          | 29           | 29           | 32          | 32           | 32         | 32       | 32         | 32         | Score          |
| 87.5       | •         | 87.5         |              | •      |              | •          | 87.5         | 87.5         | 87.5         | •                 | 87.5             | •            | •          | 90.6       | 90.6   | 90.6         | 90.6               | 90.6       | 90.6        | 90.6        | 90.6         | 90.6         | •           | •            | 100.0      | 8        | 100.0      | 8          | Query<br>Match |
| 327        | 321       | 321          | 311          | 298    | 298          | 279        | 273          | 266          | 266          | 149               | 80               | 1266         | 1266       | 1266       | 1016   | 986          | 825                | 824        | 495         | 301         | 286          | 204          | 984         | 473          | 465        | 319      | 101        | 93         | Length         |
| N          | N         | N            | N            | N      | _            | N          | N            | N            | N            | ນ                 | N                | N            |            |            |        |              | ນ                  | N          | <u>, .</u>  | N           | N            | ຎ            | N           | N            | N          | ึง       | ນ          | N          | BG             |
| AC1615     | AG3086    | B98200       | S18740       | A28260 | WMLJLK       | G97143     | H70849       | B97983       | A95114       | AE0812            | PH1166           | F91143       | A85989     | AF0911     | T25433 | G65116       | A59296             | T36818     | A26396      | AH3431      | A84797       | T04491       | AE0290      | D85041       | F71213     | A69675   | I37262     | A37263     | ID             |
| giycine be | (D        | hypothetical | bell protein | _=     | bel-1 prot   | probable n | hypothetical | hypothetical | hypothetical | conserved hypothe | Ig heavy chain V | hypothetical | hypothetic | probable e | ~      | hypothetical | alpha-L-arabinofur | probable s | T-cell sur  |             | hypothetical | hypothetical | insecticial | hypothetical | probable H | phoapho  | Ig heavy c | Ig heavy c | Description    |
| betaine/ca | ranydroio | cal prote    |              | pro    | protein - 81 |            | cal prote    |              |              | hypothet          | in V             | cal prote    |            | ~          |        |              | cabinofur          | secreted   | surface gly | zinc-cadmiu | al prote     |              |             | al prote     |            | ructokin | chain V r  | chain V r  | ğ              |

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RESULT 1
A37263
A37263
Ig heavy chain V region (4A9) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Species: Mus musculus (house mouse)
C;Accession: A37263
C;Accession: A37263
R;Goshorn, S.C.; Retzel, E.; Jemmerson, R.
J. Biol. Chem. 266, 2134-2142, 1991
A;Title: Common structural features among monoclonal antibodies binding the same antigeni A;Accession: A37263
A;Accession: A37263
   R;Goshorn, S.C.; Retzel, E.; Jemmerson, R.
J. Biol. Chem. 266, 2134-2142, 1991
J. Biol. Chem. 266, 2134-2142, 1991
A;Title: Common structural features among monoclonal antibodies binding the same antigeni A;Reference number: A38601; MUID:91115823; PMID:1703527
A;Accession: 137262
A;Status: preliminary; not compared with condeptual translation
  Ig heavy chain V region (1G1) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 30-Aug-1991 #sequence_revision 30-Aug-1991 #text_change 23-Jul-1999
C;Accession: I37262
   RESULT 2
137262
  A;Status: preliminary
A;Molecule type: mRNA
A;Molecule type: mRNA
A;Residues: 1-93 <GOS>
A;Cross-references: UNIPARC:UPI0000176AE0; GB:M57996
A;Cross-references: UNIPARC:UPI0000176AE0; GB:M57996
A;Cross-references: UNIPARC:UPI0000176AE0; GB:M57996
C;Superfamily: immunoglabulin V region; immunoglabulin homology
C;Keywords: heterotetramer; immunoglabulin
  ঠ
   A;Molecule type: mRNA
A;Residues: 1-101 <GOS>
A;Cross-references: UNIPARC:UPI0000115185; GB:M57995; NID:g195375; PIDN:AAA63334.1; PID:g
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
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  Query Match
Best Local S:
Matches 5
   Query Match
Best Local Similarity
Matches 5; Conserv
  y Match 100.0%; Score 32; DB 2; Local Similarity 100.0%; Pred. No. 3.5; hes 5; Conservative 0; Mismatches 0
  15 DYDMH 19
1 DYDMH 5
  1 DYDMH 5
  100.0%; Score 32; DB 2; Length 101; ilarity 100.0%; Pred. No. 3.9; Conservative 0; Mismatches 0; Indels
  ALIGNMENTS
  0; Indels
   Length 93;
  ç
   0
   AGA for residue 39
   Gaps
   Gaps
   0,
   0;
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RESULT 3
A69675
6-phosphofructokinase pfk - Bacillus subtilis
6-phosphofructokinase pfk - Bacillus subtilis
6-phosphofructokinase pfk - Bacillus subtilis
C;Species: Bacillus subtilis
C;Species: Bacillus subtilis
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C;Species: Bacillus subtilis
R;Sunst, F.; Ogasawara, A.; Bitan, M.; Bijtan, A.; Alloni, G.; Kuman, K.D.; Errington, J.; Fabret, C.; Ferrari, E.
R;Authors: Lauber, J.; Lazaravic, V.; Lee, S.M.; Levine, A.; Liu, H.; Masuda, S.; Hallo, M.;
R;Beger, M.; Rivolta, C.; Rocha, B.; Roche, B.; Rose, M.; Sadaie, Y.; Sato, T.; Scanlon, A;Authors: Schleich, S.; Schroeter, R.; Scoffone, F.; Sekiguchi, J.; Sekowaka, A.; Serot akeuchi, M.; Tamakoshi, A.; Tanaka, T.; Terpstra, P.; Tognoni, A.; Tosato, V.; Uchiyama, T.; Winters, P.; Mipat, A.; Zamamoto, H.; Yamane, K.; Yasumoto, K.; Yata, K.; Yoshida, K.; A; Coshikawa, H.; Danchin, A.
A;Tathus: The complete genome sequence of the Gram-positive bacterium Bacillus subtilis.
A;Status: The complete genome sequence of the Gram-positive bacterium Bacillus subtilis.
  A/Gene: PH1974

C;Superfamily: H(+)-transporting ATP synthase F;183-354/Domain: H+-transporting ATP synthase
   R;Kawarabayasi, Y.; Sawada, M.; Horikawa, H.; Haikawa, Y.; Hino, Y.; Yamamoto, S.; Seki M.; Ohfuku, Y.; Funahashi, T.; Tanaka, T.; Kudoh, Y.; Yamazaki, J.; Kushida, N.; Oguch DNA Res. 5, 55-76, 1998
A;Title: Complete sequence and gene organization of the genome of a hyper-thermophilic A;Reference number: A71000; MUID:98344137; PMID:9679194
A;Accession: F71213
   A;Gene: pfk
C;Superfamily: ATP-dependent phosphofructokinase, prokaryotic type; 6-phosphofructokina
C;4-278/Domain: 6-phosphofructokinase 1 homology <6FFl>
   A;Residues: 1-319 <KUN>
A;Cross-references: UNIPROT:O34529; UNIPARC:UPI0000608C0;
A;Cross-references: Strain 168
   A;Cross-references: UNIPROT:057729; UNIPARC:UPI0000138167; GB:AP0000007; NID:g3236134; PIA;Experimental source: strain OT3
A;Note: this accession replaces an interim accession for a sequence replaced by GenBank C;Genetics:
   probable H(+)-transporting ATP synthase subunit B - Pyrococcus horikoshii
C;Species: Pyrococcus horikoshii
C;Date: 14-Aug-1998 #sequence_revision 14-Aug-1998 #text_change 31-Dec-2004
C;Accession: F71213
  유
  8
   A;Status: preliminary; nucleic acid sequence not shown; translation not
   Query Match
Best Local
   Residues: 1-465 <KAW>
  Status: preliminary; nucleic acid sequence not shown; translation not
  Molecule type: DNA
  Local Similarity hes 5; Conserv
   Local Similarity
les 5; Conserv
  195 DYDMH 199
   1 DYDMH 5
  HWOYD
   DNA
  Conservative
   100.0%; Score 32; DB
100.0%; Pred. No. 20;
   100.0%;
  100.0%; Score 32; DB
100.0%; Pred. No. 14;
  0,
  ATP synthase alpha chain homology
  Mismatches
  DB 2; Length 319
   DB 2;
  0
   Length 465
  Indels
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   0
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  Oguchi
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  RESULT
AE0290
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A;Accession: T04491
A;Molecule type: DNA
A;Residues: 1-204 <BEV>
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RESULT 7
   insecticial toxin [imported] - Yersinia pestis (strain CO92)
(;Species: Yersinia pestis
(;Species: Yersinia pestis
(;Date: 02-Nov-2001 #sequence_revision 02-Nov-2001 #text_change 09-Jul-2004
(;Accession: AE0290
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(;Accession: AE
   R;anonymous, The European Union Arabidopsis Genome Sequencing Consortium, The Col
Nature 402, 769-777, 1999

A;Title: Sequence and analysis of chromosome 4 of the plant Arabidopsis thaliana.

A;Reference number: A85001; MUID:20083488; PMID:10617198

A;Accession: D85041

A;Status: preliminary

A;Molecule type: DNA

A;Residues: 1-473 <STO>

A;Cross-references: UNIPROT:Q9ZR05; UNIPARC:UPI00000488F1; GB:NC_001268; NID:g727

A;Gene: A74g03260

A;Map position: 4
   hypothetical protein AT4903260 [imported] - Arabidopsis thaliana
C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Date: 16-Feb-2001 #sequence_revision 16-Feb-2001 #text_change 09-Jul-2004
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A;Gene: sepC
  A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-984 <KUR>
  A; Title: Genome sequence of Yersinia pestis, the causative A; Reference number: AB0001; MUID:21470413; PMID:11586360
  A; Accession: AE0290
  A; Reference number: AB0001;
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Best Local S
Matches 5
  Matches
   Query Match
Best Local Similarity
  241
   834 DYDMH
  62 DYDMH
  1 DYDMH 5
   HWGYG 1
  Similarity
  HWOYD
  Conservative
   838
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   UNIPROT: Q9ZR05; UNIPARC: UPI00000488F1; GB: NC_001268; NID: g7270196; PJ
  100.0%;
   100.0%; Score 32; DI
100.0%; Pred. No. 21.
:ive 0; Mismatches
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  Score 32;
Pred. No.
   DB
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  0
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  agent of plague
  0
   0
  Gaps
   Gaps
  The Cold Spring
  0
   0
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O., Hoheisel,

Mewes

A; Cross references: UNII
A; Experimental source: c
C; Genetics:
A; Map position: 4
A; Note: P8P16.60

UNIPROT:049574; UNIPARC:UPI00000A96C5; EMBL:AL021633

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279 DYDLH 283

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R;Dellecchio, V.G.; Kapatral, V.; Redkar, R.J.; Patra, G.; Mujer, C.; Los, T.; Ivanova, .; Mazur, M.; Goltsman, E.; Selkov, E.; Elzer, P.H.; Hagius, S.; O'Callaghan, D.; Letess Proc. Natl. Acad. Sci. U.S.A. 99, 443-448, 2002
A;Title: The genome sequence of the facultative intracellular pathogen Brucella melitens A;Reference number: AD3252; PMID:11756688
  A; Gene: At2g37780
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   A;Title: Sequence and analysis of chromosome 2 of the plant Arabidopsis thaliana A;Reference number: A84420; MUID:20083487; PMID:10617197 A;Accession: A84797
  R;Lin, X.; Kaul, S.; Rounsley, S.D.; Shea, T.P.; Benito, M.I.; Town, C.D.; Fujii, C.Y. M.; Koo, H.; Moffat, K.S.; Cronin, L.A.; Shen, M.; VanAken, S.B.; Umayam, L.; Tallon, Jeuss, D.; Nierman, W.C.; White, O.; Eisen, J.A.; Salzberg, S.L.; Fraser, C.M.; Venter, Nature 402, 761-768, 1999
  hypothetical protein At2g37780 [imported] - Arabidopsis thaliana C;Species: Arabidopsis thaliana (mouse-ear cress) C;Date: 02-Feb-2001 #sequence_revision 02-Feb-2001 #text_change 09-Jul-2004 C;Accession: A84797
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  A; Residues: 1-286 <STO>
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8
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  0
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634 DYDLH 638

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   A;Title: Isolation of complementary DNA clones encoding the human lymphocyte glycoprotein A;Reference number: A26396; MUID:87014786; PMID:3093892 A;Accession: A26396.
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  R;Oliver, K.; Harris, D.; Parkhill, J.; Barréll, B.G.; Rajandream, submitted to the EMBL Data Library, September 1998
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   A;Cross-references: GDB:120579; OMIM:153340
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   A; Molecule type: mRNA
   Nature 323, 346-349,
S
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R;Blattner, F.R.; Plunkett III, G.; Bloch, C.A.; Perna, N.T.; E
A; Rose, D.J.; Mau, B.; Shao, Y.
Science 277, 1453-1462, 1997
A;Title: The complete genome sequence of Escherichia coli K-12.
A;Reference number: A64720; MUID:97426617; PMID:9278503
A;Accession: G65116
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submitted to the EMBL Data Library, June 1996
A;Reference number: Z20435
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A;Reference number: Z20033
A;Accession: T25433
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  Nature 413, 848-852, 2001
A;Authors: Parry, C.; Quail, M.; Rutherford, K.; Simmonds, M.;
A;Title: Complete genome sequence of a multiple drug resistant
A;Reference number: AB0502; MUID:21534947; PMID:11677608
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Result No.

| RESULT 2 Q4GYZ5_9 ID Q4G AC Q4G AC 13- DT 13- DT 13- DT Call GN ORF OS Try OC Buk                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             | рь<br>Оу                          | RP NUCLI RC STRAI RA Herri RI Submi DR EMBL SQ SEQUI Query Ma                                                                                                                                     | RP NUCLE RC STRAI RA Bart- RA Bart- RA Bray- RA Bray- RA Reitt RA Reitt RA Raree RA RA Barree RA Barree RA Barree RA Wonte                                                                                                                                                                                                                 | Q4GYZ8 9TI Q4GYZ8 9TI ID Q4GY; AC Q4GY; DT 13-S; DT 13-S; DT Calp; GN ORFN OS Tryp; OC Buka OX NCBI |            |                                                                                                                                                                                                                                                              |
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| ae; Trypanosoma.                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              | ,                                 | es.<br>;<br>h 123;<br>els 0; Gaps 0;                                                                                                                                                              | Hertz-Fowler C., arron A.J., Bowman S., C.H., Cronin A., I.S., Harper A.D., A., Rabbinowitch E., hownkeen R., Tait A., Gull K., trypanosome: gene and polymorphism.";                                                                                                                                                                      | ae; Trypanosoma.                                                                                    |            | Q9xym9 trichomonas Q531f4 oryza sativ Q4g346 emiliania h Q7qrx8 giardia lam Q5xd57 streptococc Q7cnf3 streptococc Q7cnf3 streptococc Q7xjp2 arabidopsis Q8g236 brucella su Q4psa6 arabidopsis Q8g1t2 brucella me Q8yft2 prythrobact                          |

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RA Diaz J.S., Dodge S., Ferreira P., FitzGerald M., Gage D., Galagan J.,
RA Gardyna S., Gnerre S., Graham L., Grand-Pierre N., Hafez N.,
RA Gardyna S., Gnerre S., Graham L., Grand-Pierre N., Hafez N.,
RA Hagopian D., Hagos B., Hall J., Horton L., Hulme W., Iliev I.,
RA Jaffe D., Johnson R., Jones C., Kamal M., Kamat A., Karatas A.,
Ra L.-J., Mabbitt R., MacLean C., Macdonald P., Major J., Manning J.,
Ma L.-J., Mabbitt R., MacLean C., Macdonald P., Major J., Manning J.,
Mathews C., Muzphy T., Naylor J., Nguyen C., Nicol R.,
RA Milova T., Mlenga V., Muzphy T., Naylor J., Nguyen C., Nicol R.,
RA Nielsen C.B., Norbu C., O'Connor T., O'Donnell P., O'Neil D.,
Nielsen C.B., Norbu C., O'Connor T., O'Donnell P., O'Neil D.,
RA Nielsen C., Ramasamy U., Raymond C., Retta R., Rise C., Rogov P.,
RA Rachupka A., Ramasamy U., Raymond C., Retta R., Rise C., Rogov P.,
RA Rachupka A., Ramasamy U., Raymond C., Retta R., Rise C., Rogov P.,
RA Ramas J., Teafaye S., Theodore J., Topham K., Travers M.,
Talamas J., Teafaye S., Theodore J., Topham K., Travers M.,
Na Talamas J., Yenkataraman V.S., Viel R., Vo A., Wang S., Wilson B.,
Na M. Y., Wang D., Young G., Zainoun J., Zembek L., Zimmer A., Zody M.,
  Query Match
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Matches 5
  NUCLEOTIDE SEQUENCE.
STRAIN=FGSC A4,
Birren B., Nusbaum C.,
Arachchi H.M., Barna I
  STRAIN=927/4 GUTat10.1;
Hertz-Fowler C., Berriman M.;
Hertz-Fowler C., Berriman M.;
Submitted (JUN-2005) to the EMBL/GenBank/DDBJ databases.
EMBL; AL929603; CAJ16336.1; -; Genomic_DNA.
SEQUENCE 129 AA; 14030 MW; 91C38ECF31609A1A CRC64;
   Hall N., Berriman M., Lennard N.J., Harris B.R., Hertz-Fowler C.,
Bart-Delabesse B.N., Gerrare C.S., Atkin R.J., Barron A.J., Bowman S
Bray-Allen S.P., Bringaud F., Clark L.N., Corton C.H., Cronin A.,
Davies R., Doggett J., Fraser A., Gruter B., Hall S., Harper A.D.,
Kay M.P., Leech V., Mayes R., Price C., Quail M.A., Rabbinowitch B.,
Kay M.P., Leech V., Mayes R., Price C., Quail M.A., Rabbinowitch B.,
Reitter C., Rutherford K., Sasse J., Sharp S., Shownkeen R.,
Racled A., Taylor S., Tweedie A., Turner C.M.R., Tait A., Gull K.,
Barrell B., Melville S.E.,
The DNA sequence of chromosome I of an African trypanosome: gene
content, chromosome organisation, recombination and polymorphism.",
   10-MAY-2005 (TrEMBLrel. 30, 10-MAY-2005 (TrEMBLrel. 30, 10-MAY-2005 (TrEMBLrel. 30, 10-MAY-2005 (TrEMBLrel. 30, Hypothetical protein. ORFNames=AN5781.2;
   Aspergillus nidulans FGSC A4.
Eukaryota; Fungi; Ascomycota; Pezizomycotina; Eurotiomycetes;
Eurotiales; Trichocomaceae; Emericella.
  Submitted
   Q5B0Z9 EMENI PRELIMINARY;
   STRAIN=927/4 GUTat10.1;
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  "Genome Sequence of Aspergillus nidulans.";
Submitted (JAN-2004) to the EMBL/GenBank/DDBJ
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   Acids Res. 31:4864-4873 (2003).
   DYDMH 58
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   100.0%; Score 32; DB 2; Length 129; ilarity 100.0%; Pred. No. 47; Conservative 0; Mismatches 0: Trdala
   Created)
Last sequence update)
Last annotation update)
                         Bank/DDBJ databases
is derived from an
   276
   8
   Galagan J.,
  0;
   Bowman S.,
   Gaps
   0
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ID AC LE
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  RESULT 5
K6PF_BACSU
   문
   Query Match
Best Local S
Matches 5
   Query Match
Best Local S
Matches 5
15-JUL-1998 (Rel. 36, C
15-JUL-1998 (Rel. 36, I
13-SEP-2005 (Rel. 48, I
6-phosphofructokinase (
   EMBL; AACDO10
Hypothetical
SEQUENCE 27
   "The complete genome sequence of Francisella tularensis, the causative agent of tularenda.";
Nat. Genet. 37:153-159(2005).

EMBL; AJ749949; CR645244.1; -; Genomic DNA.
GO; GO:0008800; F:beta-lactamase activity; IEA.
GO; GO:0016787; F:hydrolase activity; IEA.
GO; GO:0016787; P:beta-lactam antibiotic catabolism; IEA.
GO; GO:003655; P:beta-lactam antibiotic; IEA.
InterPro; IPR0001466; Beta_lactamase_A.
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InterPro; IPR000008; C2.
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   Larsson P., Oyston P.C.F., Chain P., Chu M.C., Duffield M., Fuxelius H.-H., Garcia E., Haelltorp G., Johansson D., Isherwoo, Karp P.D., Larsson E., Liu Y., Michell S., Prior J., Prior R., Malfatti S., Sjoestedt A., Svensson K., Thompson N., Vergez L., Wagg J.K., Wren B.W., Lindler L.E., Andersson S.G.E., Forsman M. Titball R.W.,
  01-FEB-2005 (TrEMBLrel. 29, I
01-FEB-2005 (TrEMBLrel. 29, I
Beta-lactamase (EC 3.5.2.6).
  PRINTS; PRO0118; BLACTAMASEA.
PRINTS; PRO0360; C2DOWAIN.
PROSITE; PS00146; BERG LACTAMASE A;
  STRAIN=SCHU S4 / Schu 4;
PubMed=15640799; DOI=10.1038/ng1499;
  Francisella tularensis (subsp. tularensis).
Bacteria; Proteobacteria; Gammaproteobacteria; Thiotrichales;
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  NCBI_TaxID=119856;
  Francisellaceae; Francisella.
   01-FEB-2005
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L; AACDD1000098; EAA62874.1; -; Genomic_DNA.
   PF00144; Beta-lactamase;
   73
  92 DYDMH 96
  1 DYDMH 5
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   Similarity
5; Conserv
   1 DYDMH
  ū
  Similarity
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287 AA; 31982 MW; 93FC6A24A1C0383A CRC64;
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  Conservative
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(Rel. 36, Last seq
(Rel. 48, Last ann
  protein
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   31803 MW;
  100.0%;
   100.0%;
 Last sequence update)
Last annotation update)
(EC 2.7.1.11) (Phosphofructokinase)
  Last sequence update)
Last annotation updat
  Created)
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  0
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Pred. No. 1.1
D; Mismatches
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Pred. No. 1.1e+02;
   9C1E24C76A107E74
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   DB 2;
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  RA Kunst F. Ogasawara N., Moszer I., Albertini A.M., Alloni G.,
RA Azevedo V., Bertero M.G., Bessieres P., Bolotin A., Borchert S.,
RA Borriss R., Boursier I., Brans A., Braun M., Brignell S.C., Bron S.,
RA Brouillet S., Bruschi C.V., Caldwell B., Capuano V., Carter N.M.,
RA Choi S.-K., Codani J.-J., Connerton I.F., Cummings N.J., Daniel R.A.,
RA Choi S.-K., Codani J.-J., Fabret C., Ferrari B., Follger D.,
RA Denizot F., Devine K.M., Duesterhoeft A., Ehrlich S.D., Emmerson P.T.,
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RA Guiseppi G., Guy B.J., Haga K., Halech J., Harwood C.R., Henaut A.,
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RA Jones L.-M., Joris B., Karamate D., Kasahara Y., Klaerr-Blanchard M.,
RA Medigue C., Medina N., Mellado R.P., Mizuno M., Moest J.D., Nakai S.,
RA Kumano M., Kurita K., Lapidus A., Lardinois S., Lauber J.,
RA Priscott J., Persecan B., Pujic P., Purnelle B., Rapoport G.,
RA Prescott A.M., Presecan B., Pujic P., Purnelle B., Rapoport G.,
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RA Rose M., Sadaie Y., Sato T., Sekwaka N., Seror S.J., Serror P.,
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RA Takemaru K., Takeuchi M., Tamakoshi A., Tanaka T., Takadi T., Takahashi H.,
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RA Vassarotti A., Wandri A., Wandenbol M., Yamane K.,
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   HSSP; Put-
MR; 034529;
  Lapidus A., Galleron N., Sorokin A., I 
"Sequencing and functional annotation 
in the 200 kb rrnB-dnaB region.";
Subtilist; EG12644; pfkA.

HAMAP; MF 00339; -; 1.

InterPro; IPR012003; ATP PFK prok.

InterPro; IPR012028; PFKA ATP.

InterPro; IPR000023; Ppfruckinase.

PANTHER; PTHR13697; Ppfruckinase; 1.
   MEDLINE=98044033; PubMed=9384377; DOI=10.1038/36786;
  Microbiology 143:3431-3441(1997)
  (Phosphohexokinase).
Name=pfkA; Synonyms=pfk; OrderedLocusNames=BSU29190;
Bactllus subtills.
Bacteria; Firmicutes; Bacillales; Bacillaceae; Bacillus.
  EMBL; AF008220; AAC00342.1; -; GEMBL; Z99118; CAB14879.1; -; GEPIR; A69675, A69675.
  This Swiss-Prot entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation - the European Bioinformatics Institute. There are no restrictions on its use as long as its content is in no way modified and this statement is not
   "The complete genome sequence of the Gram-positive bacterium Bacillus subtilis.";
   MEDLINE=98048467; PubMed=9387221;
  removed.
   Nature 390:249-256(1997).
  Yoshikawa H., Danchin
  CATALYTIC ACTIVITY: ATP + D-fructose 6-phosphate =
   SEQUENCE [LARGE SCALE
  1-319
  Genomic_DNA
   Genomic_
   GENOMIC
   Ehrlich
  Ehrlich S.-D.;
of the Bacillus subtilis genes
  DNA].
   ADP
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   SOUTH FILE STATES OF THE STATE
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RA Arachchi H.M., Barna N., Bastlen V., Bløm T., Boguslavkiy L.,
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RA Arachugka A., Ramasamy U., Raymond C., Reitta R., Rise C., Rogov P.,
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13-SEP-2005
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NP_BIND
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PROSITE; PS00433; PHOSPHOFRUCTOKINASE; 1.

Allosteric enzyme; ATP-binding; Complete pallosteric enzyme; ATP-binding; Notide-binding; Noticetide-binding;   Pfam; PF00365; PFK; 1.

PIRSF; PIRSF000532; ATP PFK prok; ?

PRINTS; PR00476; PHFRCTKINASE.

PRODOM; PD000707; Ppfruckinase; 1.
   Q4IHZ6;
                      Wu X., "YHELL. HEADOR S., "FLEATHER E.;
"FLEAR'S (PEB-2004) to the EMBL/GenBank/DDBJ databases.
Submitted (PEB-2004) to the EMBL/GenBank/DDBJ databases.

TATION: The sequence shown here is derived from an
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ORFNames=FG03162.1;
   SEQUENCE
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   Hypocreomycetidae; Hypocreales; NCBI_TaxID=229533;
  Eukaryota; Fungi; Ascomycota;
  Gibberella zeae PH-1.
   STRAIN-PH-1
  NUCLEOTIDE SEQUENCE.
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  1 DYDMH 5
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   319 AA;
   (TrEMBLrel. 31, Created)
(TrEMBLrel. 31, Last sequence update)
(TrEMBLrel. 31, Last annotation update)
   Conservative
   187
162
243
249
252
  25
158
187
187
127
   100.0%;
   ₩;
  ATP (By similarity).

ATP (By similarity).

ATP (By similarity).

ATP (By similarity).

Proton acceptor (By similarity).

Magnesium (via carbonyl oxygen) (
similarity).

Magnesium (By similarity).

Substrate (By similarity).
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   Substrate
   Substrate
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Pred. No.
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   326 AA.
   (By similarity).
(By similarity).
(By similarity).
(By similarity).
(By similarity).
  1.3e+02;
hes 0;
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   (By
  Gaps
  0
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Q7N656
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ID Q7N656
ID Q7N656
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   Query Match
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Matches 5
   Query Match
Best Local Similarity
Matches 5; Conserv
   Q8U4A5;
28-FEB-2003 (Rel. 41, Created)
28-FEB-2003 (Rel. 41, Last sequence update)
10-MAY-2005 (Rel. 47, Last annotation update)
10-MAY-2005 (Rel. 47, Last annotation update)
    NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC DNA].
STRAIN=Vc1 / DSM 3638 / ATCC 43587 / JCM 8422;
Weiss R.B., Dunn D.M., Robb F.T., Brown J.R.;
"The complete sequence of the Pyrococcus furiosus genome.";
Submitted (FEB-2002) to the EMBL/denBank/DDBJ databases.
-i- FUNCTION: Produces ATP from ADP in the presence of a proton
  PYRFU
   MEDLINE-22957627; PubMed=14528314; DOI=10.1038/nbt886;
Duchaud E., Rusniok C., Frangeul L., Buchrieser C., Givaudan A.,
Taourit S., Bocs S., Boursaux-Eude C., Chandler M., Charles J.-F.,
Dassa E., Derose R., Derzelle S., Freyssinet G., Gaudriault S.,
Medigue C., Lanois A., Powell K., Siguier P., Vincent R., Wingate V.
Zouine M., Glaser P., Boemare N., Danchin A., Kunst F.;
"The genome sequence of the entomopathogenic bacterium Photorhabdus luminescens.";
  01-MAR-2004 (TrEMBLrel. 26, Created)
01-MAR-2004 (TrEMBLrel. 26, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
01-MAR-0004 (TrEMBLrel. 26, Last annotation update)
Similarities with C-terminal region of insecticial toxin.
OrderediocusNameseplu1712;
Photorhabdus luminescens (subsp. laumondii).
Bacteria, Proteobacteria, Gammaproteobacteria, Enterobacteriales;
  EMBL; AACM01000148;
Hypothetical proteir
SEQUENCE 326 AA;
   Q7N656 PHOLL PRELIMINARY; Q7N656;
   Pyrococcus furiosus.
Archaea, Euryarchaeota,
  SEQUENCE
   Nat. Biotechnol. 21:1307-1313(2003).
EMBL; BX571864; CAE14005.1; -; Genomic_DNA.
PhotoList; plu1712; -.
Complete proteome.
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  Enterobacteriaceae; Photorhabdus.
  NCBI_TaxID=2261;
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   PYRFU
  292
  194 DYDMH 198
  1 DYDMH
  1 DYDMH 5
   Similarity
5; Conserv
  DYDMH 296
   340 AA;
   Conservative
   Conservative
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Pred. No. 1.4e+02;
   PRT;
   F211C78983428339 CRC64,
  Mismatches
  Mismatches
  Thermococcales;
   462 AA.
  340
  ጅ
  <u>.</u>
  Length 340
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  Indels
  Thermococcaceae;
   R., Wingate V.,
  Photorhabdus
   0
   subunit
   0
   Gape
   Gaps
   0
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RESULT 9
VATB_PYRHO
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HAMAR; MF 00310; -; 1.
InterPro; IPR005724; ATP synthA B.
InterPro; IPR000793; ATPase_a/b_C.
InterPro; IPR000194; ATPase_a/b N.
InterPro; IPR000194; ATPase_a/b Centre.
Pfam; PF00006; ATP-synt_ab; 1.
Pfam; PF00306; ATP-synt_ab C; 1.
Pfam; PF00306; ATP-synt_ab C; 1.
Pfam; PF02874; ATP-synt_ab N; 1.
IGREAMS; TIGR01041; ATP syn B arch; 1.
PROSITE; PS00152; ATPASE_ALPHA_BETA; 1.
   Kawarabayasi Y., Sawada M., Horikawa H., Haikawa Y., Hino Y., Yamamoto S., Sekine M., Baba S.-I., Kosubi H., Hosoyama A., Nagai Y., Sakai M., Ogura K., Otsuka R., Nakazawa H., Takamiya M., Ohfuku Y., Funahashi T., Tanaka T., Kudoh Y., Yamazaki J., Kushida N., Oguchi A., Aoki K.-I., Yoshizawa T., Nakamura Y., Robb P.T., Horikoshi K., Masuchi Y., Shizuya H., Kikuchi H., "Complete sequence and gene organization of the genome of a hyper-thermophilic archaebacterium, Pyrococcus horikoshii OT3.", DNA Res 5:55-76(1998).

-I- FUNCTION: Produces ATP from ADP in the presence of a proton gradient across the membrane. The archaeal beta chain is a
  VATB PY:
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  STRAIN=OT3;
MEDLINE=98344137; PubMed=9679194;
  This Swiss-Prot entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use as long as its content is in no way modified and this statement is not
  Name=atpB; OrderedLocusNames=PH1974;
Pyrococcus horikoshii.
Archaea; Euryarchaeota; Thermococci;
   ATP synthesis; Complete proteome; Hydrogen ion transport; Ion transport; Transport.
SEQUENCE 462 AA; 51915 MW; 5AB9FBAFA2C86283 CRC64;
   -!- SIMILARITY: Belongs to the ATPase alpha/beta chains family.
  NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC DNA]
  removed
  NCBI_TaxID=53953;
   Pyrococcus.
  16-OCT-2001 (Rel. 40, Created)
16-OCT-2001 (Rel. 40, Last sequence update)
10-MAY-2005 (Rel. 47, Last annotation update)
7-type ATP synthase beta chain (EC 3.6.3.14)
  regulatory subunit.
CATALYTIC ACTIVITY:
   PYRHO
   238 DYDMH 242
   1 DYDMH 5
  l Similarity
5; Conserv
  100.0%; ilarity 100.0%; Conservative 0
  STANDARD;
  Thermococci; Thermococcales;
   ATP + H(2)O + H(+)(In)
   0
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Pred. No.
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  Genomic_DNA.
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  465
   1.9e+02
  B
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  (V-type ATPase subunit
   n
  Length 462
   ADP
  ADP + phosphate
   + phosphate
  Thermococcaceae;
  Hydrolase;
   <u>,</u>
   Gaps
   0
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  밁
   Query Match
Best Local
  HAMAP; MF_00310; -; 1.

InterPro; IPR005724; ATP synthA B.

InterPro; IPR00793; ATPase_a/b_C.

InterPro; IPR00194; ATPase_a/b_N.

InterPro; IPR00194; ATPase_a/bcentre.

R InterPro; IPR00194; ATPase_a/bcentre.

R Pfam; PP00306; ATP-synt_ab; 1.

R Pfam; PP00306; ATP-synt_ab_N; 1.

R PR0SITE; PS00152; ATPASE ALPHA BETA; 1.

W ATP synthesis; Complete proteome; Hydrogen ion transport; Hydroge
  Matches
   01-MAY-1999 (TrEMBLrel. 10, Created)
01-MAY-1999 (TrEMBLrel. 10, Last sequence update)
10-MAY-2005 (TrEMBLrel. 30, Last sequence update)
10-MAY-2005 (TrEMBLrel. 30, Last annotation update)
Hypothetical protein F4C21.19 (Hypothetical protein AT4g03260).
Name=F4C21.19; Synonyms=AT4g03260;
Arabidopsis thaliana (Mouse-ear cress).
Eukaryota, Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnollophyta; eudicotyledons; core eudicotyledons;
rosids; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
  IT 10
ARATH
Q5ZROS_ARATH PRELIMINARY;
   NUCLEOTIDE SEQUENCE.
Habermann K., de la Bastide M., Huang Preston R., Calma C., Martienssen R.,
  SEQUENCE
  EMBL;
  иве
   EU Arabidopsis sequencing project; submitted (MAR-2000) to the EMBL/GenBank/DDBJ EMBL; AC005275; AAD14454.1; -; Genomic_DNA. EMBL; AL161496; CAB77811.1; -; Genomic_DNA. PIR; D85041; D85041.
   Spiegel L.A., Huang E.N., Nascimento L.U., de Preston R.R., Matero A., Shah R., O'Shaughness Shekher M., Schutz K., See L.H., Swaby I., Hak Mewes H.W., Lemcke K., Mayer K.F.X.; Submitted (MAR-2000) to the EMBL/GenBank/DDBJ
   Q9ZR05
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   NUCLEOTIDE SEQUENCE
  McCombie W.R.;
"Arabidopsis thaliana
  NCBI_TaxID=3702;
PRINTS; PR00019; LEURICHRPT.

Hypothetical protein; Leucine-rich repeat; Repeat.

SEQUENCE 473 AA; 52344 MW; 02EF2385906D5C40 CRC64;
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InterPro; IPR007092; LRR_SDS22.
Pfam; PF00560; LRR_1; 4.
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  241 DYDMH 245
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   l Similarity
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  1 DYDMH 5
  (FEB-1999)
   (JUL-1998)
   rt; Transport.
465 AA; 52265 MW; EDAFDEF4D74C2A54 CRC64;
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  BAA31101.1; -;
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  Score 32; DB 1;
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   IV near 17 cM.";
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InterPro; IPR003056; Ig-MHC.
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SEQUENCE 493 AA; 53224 MW; 12ECD7E
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25-OCT-2004 (*25-OCT-2004 (*
   TISSUE-Mammary gland;
Otsuki T., Ota T., Nishikawa T., Hayashi K., Suzuki Y., Yamamoto J.,
Wakamatsu A., Kimura K., Sakamoto K., Hatano N., Kawai Y., Ishii S.,
Saito K., Kojima S., Sugiyama T., Ono T., Okano K., Yoshikawa Y.,
Saito K., Sasaki N., Hattori A., Okumura K., Nagai K., Sugano S.,
   01-OCT-2002 (TrEMBLrel. 22, Created)
01-OCT-2002 (TrEMBLrel. 22, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
CDNA FLJ90170 fis, clone MAMMA1000370, highly similar to Ig alpha-1
  25-OCT-2004 (TrEMBLrel. 28, 25-OCT-2004 (TrEMBLrel. 28, 25-OCT-2004 (TrEMBLrel. 28, 47)pothetical protein. ORFNames=GZ27A8_54;
  Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
  QBNCL6;
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   Isogai T.;
"Signal Sequence and Keyword Trap in silico
rength Human cDNAs Encoding Secretion or Men
Capped cDNA Libraries.";
DNA Res. 12:117-126(2005).
  NCBI_TaxID=9606;
  chain C region
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Archaea; environmental sample
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   MEDLINE=22386257; PubMed=12477932; DOI=10.1073/pnas.242603899;

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X Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

XA Alischul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

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XA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

XA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

XA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

XA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

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XA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

XA Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratte P.H.,

XA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

XA Willalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

XA Willalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

XA Pahey J., Helton B., Ketteman M., Madan A., Rodrigues S., Sanchez A.,

XA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

XA Butterfield Y.S.N., Xrzywinski M.I., Skalska U., Smailus D.E.,

XA Botherch A., Schein J.E., Jones S.J.M., Marra M.A.,

**Generation and initial analysis of more than 15,000 full-length human

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  Q6P878 XENTR
Q6P878;
05-JUL-2004 (
  PubMed=15353801; DOI=10.1126/science.1100025; Hallam S.J., Putnam N., Preston C.M., Detter J.C., Rokhsar D., Richardson P.M., DeLong B.F.; Richardson P.M., DeLong B.F.; Reverse methanogenesis: testing the hypothesis with environme: genomics.";
  Klein S., Gerhard D.S.;
Submitted (NOV-2003) to the EMBL/GenBank/DDBJ databases
EMBL; BC061351; AAH61351.1; -; mRNA.
Hypothetical protein.
SEQUENCE 562 AA; 66612 MW; C0393E711095E3D1 CRC64;
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Xenopodinae; Xenopus; Silurana.
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  05-JUL-2004 (TrEMBLrel. 27, Created)
05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
Hypothetical protein MGC75896.
   Putnam N., Detter J.C., Richardson P.M., Rokhsar D.; Submitted (AUG-2004) to the EMBL/GenBank/DDBJ databases. EMBL; AY714844; AAU83234.1; -; Genomic_DNA. Hypothetical protein.
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Pred. No.
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  EMBL; AF077207; AAD27002.1;
EMBL; AF083243; AAD39841.1;
EMBL; BC001101; AAH01101.1;
EMBL; BC007510; AAH07510.1;
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   NCBI_TaxID=9606;
   Name=EIF3S6IP; ORFNames=HSPC021, HSPC025;
  HUMAN
      HGNC:18138; BIF3S6IP
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   Pong
   Res. 10:1546-1560(2000).
  HWGYG
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TISSUE-Brain, Cervix, and Colon;

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RA Strausberg R.L., Feingold B.A., Grouse L.H., Derge J.G.,

RA Klausner R.D., Collins F.S., Wagner L., Bhenmen C.M., Schuler G.D.,

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RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsleh F.,

RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

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RA Rosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,

RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulhy S.W.,

RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulhy S.W.,

RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulhy S.W.,

RA Fahey J., Helton E., Ketteman M., Madan Å., Rodrigues S., Sanchez A.,

RA Fahey J., Helton E., Ketteman M., Madan Å., Rodrigues S., Sanchez A.,

RA Fahes J., Hulwa M., Green E.D., Dickson M.C.,

RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,

RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,

"Generation and initial analysis of more than 15,000 full-length human

RT and mouse cDNA sequences.";

RT Janches S. J.M., Marra M.A.;

RT Janches S. J. Marla M.A., Janches S. J.M., Marra M.A.;

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  Q9Y262;
28-FEB-2003 (Rel. 41, Created)
28-FEB-2003 (Rel. 41, Last sequence update)
13-SEP-2005 (Rel. 48, Last annotation update)
Eukaryotic translation initiation factor 3 subunit
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TISSUB-Umbilical cord blood;

MEDLINE=20499367; PubMed=11042152; DOI=10.1101/gr.140200;

Zhang Q.-H., Ye M., Wu X.-Y., Ren S.-X., Zhao M., Zhao C.-J., Fu G.,

Zhang Q.-H., Ye M., Wu X.-Y., Ren S.-X., Enao M., Zhao G., Zhang J.-W.,

Shen Y., Fan H.-Y., Lu G., Zhong M., Xu X.-R., Han Z.-G., Zhang J.-W.,

Tao J., Huang Q.-H., Zhou J., Hu G.-X., Gu J., Chen S.-J., Chen S.,

"Cloning and functional analysis of cDNAs with open reading frames for 300 previously undefined genes expressed in CD34+ hematopoietic
  This Swiss-Prot entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the Eventhe European Bioinformatics Institute. There are no restr
   :. Natl. Acad. Sci. U.S.A. 99:16899-16903 (2002)
  Q9H0D6:XRN2; NbExp=1; IntAct=EBI-373519, EBI-372110;
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01-FEB-2005 (TrEMBLrel. 29, Last annotation update)
   InterPro; IPR011990; TPR-like_helical. Protein biosynthesis. SEQUENCE 564 AA; 66727 MW; ECBD321
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Hui R.T., Liu Y.Q., Liu B., Zhao B., Meng X.M., Sheng H., Xu Y.Y.,
Wang X.Y., Ye J., Song L., Gao Y., Wei Y.J., Zhang C.L., Zhang J.,
Chai M.Q., Chen J.Z., Sun Y.H., Zhou X.L., Jiang Y.X., Zhao X.W.,
Liu S., Cao H.Q., Zhao Y., Liu D.Q., Ding J.F., Liu L.S., Gao R.L.,
Mu Q.Y., Qiang B.Q., Yuan J.G., Liew C.C., Zhao M.S.;
Submitted (NOV-1998) to the EMBL/GenBank/DDBJ databases.
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  GenCore version 5.1.6
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US-09-252-991A-25364
US-09-253-971-45590
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                      Sequence 5904, Ap Sequence 5112, Ap Sequence 5112, Ap Sequence 5113, Ap Sequence 5114, Ap Sequence 5114, Ap Sequence 5111, Ap Sequence 5111, Ap Sequence 6151, Ap Sequence 60718, A Sequence 60718, A Sequence 60718, A Sequence 45241, A Sequence 45241, A Sequence 45241, A Sequence 59924, A Sequence 5994, Appli Sequence 5208, Appli Sequence 5208, Appli Sequence 45241, A Sequence 5994, Ap Sequence 5208, Appli Sequence 5208, Appli Sequence 5208, Appli Sequence 45241, A Sequence 5208, Appli Sequence 5208, Appli Sequence 5208, Appli Sequence 5208, Appli Sequence 5208, Appli Sequence 5208, Appli Sequence 5208, Ap Sequence 5208, Ap Sequence 5208, Ap Sequence 5208, Ap Sequence 52364, A Sequence 53364, A
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| WS-09-513-999C-5024  US-09-513-999C-5024, Application US/09513999C  Patent No. 6783961  GENERAL INFORMATION: APPLICANT: Dumas Milne Edwards, J.B. APPLICANT: Dumas Milne Edwards, J.B. APPLICANT: Dumas Milne Edwards, J.B.  APPLICANT: Dumas Milne Edwards, J.B.  APPLICANT: Giordano, J.Y.  TITLE OP INVENTION: Expressed Sequence Tags and Encoded Human Proteins.  Patent No. 6783961  CURRENT APPLICATION NUMBER: US/09/513,999C  CURRENT FILING DATE: 2000-02-24  PRIOR APPLICATION NUMBER: US 60/122,487  PRIOR FILING DATE: 1999-02-26  NUMBER OF SEQ ID NOS: 36681  SOFTWARE: Patent.pm | Query Match 100.0%; Score 32; DB 2; Length 93; Best Local Similarity 100.0%; Pred. No. 16; Matches 5; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  Qy 1 DYDMH 5 Db 22 DYDMH 26 | RESULT 1 US-09-621-976-5904 ; Sequence 5904, Application US/09621976 ; Patent No. 6639063 ; Patent No. 6639063 ; GENERAL INFORMATION: APPLICANT: Dumas Milne Edwards, J.B. APPLICANT: Jobert, S. APPLICANT: Jobert, S. APPLICANT: Jobert, S. FILE OF INVENTION: ESTs and Encoded Human Proteins. FILE REFERENCE: GENSET.054PR2 CURRENT APPLICATION NUMBER: US/09/621,976 CURRENT APPLICATION NUMBER: US/09/621,976 CURRENT FILING DATE: 2000-07-21 NUMBER OF SEQ ID NOS: 19335 SOFTWARE: Patent.pm SEQ ID NO 5904 ; LENGTH: 93 TYPE: PRT ORGANISM: Homo sapiens FEATURE: NAME/KEY: SIGNAL I LOCATION: -821 US-09-621-976-5904 | 28 27 84.4 99 2 US-09-270-747-60648 Sequence 60648, A 29 27 84.4 116 1 US-07-634-278-57 Sequence 56, Appl 30 27 84.4 116 1 US-07-634-278-73 Sequence 73, Appl 31 27 84.4 116 1 US-08-477-728-57 Sequence 56, Appl 32 27 84.4 116 1 US-08-477-728-57 Sequence 56, Appl 33 27 84.4 116 1 US-08-477-728-57 Sequence 56, Appl 34 27 84.4 116 1 US-08-477-728-73 Sequence 57, Appl 35 27 84.4 116 1 US-08-474-040-57 Sequence 57, Appl 36 27 84.4 116 1 US-08-474-040-73 Sequence 57, Appl 37 27 84.4 116 1 US-08-474-040-73 Sequence 57, Appl 38 27 84.4 116 1 US-08-472-000-57 Sequence 56, Appl 40 27 84.4 116 1 US-08-487-200-57 Sequence 57, Appl 41 27 84.4 116 2 US-08-487-200-73 Sequence 57, Appl 42 27 84.4 116 2 US-08-487-200-73 Sequence 57, Appl 43 27 84.4 116 2 US-08-487-200-73 Sequence 57, Appl 44 27 84.4 116 2 US-08-484-537-57 Sequence 73, Appl 45 27 84.4 116 2 US-08-484-537-73 Sequence 73, Appl 46 27 84.4 116 2 US-08-484-537-73 Sequence 73, Appl 47 27 84.4 116 2 US-08-484-537-73 Sequence 73, Appl 48 27 84.4 116 2 US-08-484-537-73 Sequence 73, Appl 49 27 84.4 116 2 US-08-484-537-73 Sequence 73, Appl 40 27 84.4 116 2 US-08-484-537-73 Sequence 73, Appl 41 27 84.4 116 2 US-08-484-537-73 Sequence 73, Appl 42 27 84.4 116 2 US-08-484-537-73 Sequence 73, Appl 43 27 84.4 116 2 US-08-484-537-73 Sequence 73, Appl 44 27 84.4 116 2 US-08-484-537-73 Sequence 73, Appl |
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SEQ ID NO 5024

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Sequence 5116, Application US/09621976

Fatent No. 16639063

GENERAL INFORMATION:
APPLICANT: Dumas Milne Edwards, J.B.
APPLICANT: Dumas Milne Edwards, J.B.
APPLICANT: Jobert, S.
APPLICANT: Jobert, S.
APPLICANT: Jobert, S.
FILE COPINCENTION: ESTS and Encoded Human Proteins.
FILE REFERENCE: GENEST.054PR2
CURRENT APPLICATION NUMBER: US/09/621,976
CURRENT FILING DATE: 2000-07-21
INUMBER OF SEQ ID NOS: 19335
SOFTWARE: Patent.pm
SEQ ID NO 5116
LENGTH: 149
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ORGANISM: Homo sapiens
  APPLICANT: Dumas Milne Edwards, J.B.
APPLICANT: Jobert, S.
APPLICANT: Giordano, J.Y.
APPLICANT: Giordano, J.Y.
ITITLE OF INVENTION: ESTs and Encoded Human Proteins.
FILE REFERENCE: GENSET. 054PR2
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CURRENT FILING DATE: 2000-07-21
CURRENT FILING DATE: 2000-07-21
NUMBER OF SEQ ID NOS: 19335
SOFTWARE: Patent.pm
SEQ ID NO 5112
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APPLICANT: Dumas Milne Edwards, J.B.
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  APPLICANT: Jobert, S.
APPLICANT: Glordano, J.Y.
TITLE OF INVENTION: ESTE and Encoded Human Proteins.
FILE REFERENCE: GENSET.054PR2
CURRENT APPLICATION NUMBER: US/09/621,976
CURRENT FILING DATE: 2000-07-21
                         APPLICANT: Dumas Milne Edwards, J.B.
APPLICANT: Jobert, S.
APPLICANT: Giordan TV
TITLE OF INVENTION: ESTs and Encoded Human Proteins.
   CURRENT APPLICATION NUMBER: US/09/621,976
CURRENT FILING DATE: 2000-07-21
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   APPLICANT: Dumas Milne Edwards, J.B. APPLICANT: Jobert, S. APPLICANT: Giordano, J.Y.
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   APPLICANT: Jobert, S.
APPLICANT: Glordano, J.Y.
APPLICANT: Glordano, J.Y.
TITLE OF INVENTION: ESTS and Encoded Human Proteins.
FILE REFERENCE: GENSET.054PR2
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CURRENT FILING DATE: 2000-07-21
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ORGANISM: Homo sapiens
  APPLICANT: Giordano,
  APPLICANT: Dumas Milne Edwards, J.B.
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APPLICANT: GARY BRETON
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   APPLICANT: Schweizer, Johannes
APPLICANT: Arbor Vita Corporation
TITLE OP INVENTION: Molecular Interactions in Hematopoietic
TITLE OP INVENTION: Cells
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OR FILING DATE: 1999-10-21
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GENERAL INFORMATION:

APPLICANT: Goldman, Barry S.

APPLICANT: Hinkle, Gregory J.

APPLICANT: Slater, Steven C.

APPLICANT: Slater, Steven C.

APPLICANT: Wiegand, Roger C.

FILE REFERENCE: 38-10(15849)B

CURRENT APPLICATION NUMBER: US/09/902,540

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CURRENT APPLICATION NUMBER: 60/217,883

PRIOR APPLICATION NUMBER: 60/217,883

PRIOR FILING DATE: 2000-07-10

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Patent No. 6610836

GENERAL INFORMATION:
APPLICANT: GRAY BIFCON et. al
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO KLEBSIELLA
TITLE OF INVENTION: PNEUMONIAE FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 2709.2004001
CURRENT APPLICATION NUMBER: US/09/489,039A
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PRIOR APPLICATION NUMBER: US 60/117,747
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CURRENT FILING DATE: 1999-66-30
  APPLICANT, Homburger et al. TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster FILE REFERENCE: File Reference: 7326-094 CURRENT APPLICATION NUMBER: US/09/270,767
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PRIOR PILING DATE: 1997-10-31
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GenCore version 5.1.6
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  Description
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| RESULT 2 US-10-632-706-163 ; Sequence 163, Application US/10632706 ; Publication No. US20040175385A1 ; GENERAL INFORMATION: ; APPLICANT: MARKS, JAMES D. ; APPLICANT: MARKS, JAMES D. ; TITLE OF INVENTION: THERAPEUTIC MONOCLONAL ANTIBODIES THAT NEUTRALIZE BOTULINUM ; TITLE OF INVENTION: NEUROTOXINS ; FILE REFERENCE: 407T-895120US ; CURRENT APPLICATION NUMBER: US/10/632,706 ; CURRENT FILING DATE: 2003-08-01 ; PRIOR APPLICATION NUMBER: US 60/400,721 ; PRIOR APPLICATION NUMBER: US 60/400,721 ; PRIOR APPLICATION NUMBER: US 09/144,806 | Query Match 100.0%; Score 32; DB 4; Length 5; Best Local Similarity 100.0%; Pred. No. 1.7e+06; Matches 5; Conservative 0; Mismatches 0; Indels 0; Gaps 0; Qy 1 DYDMH 5 | RESULT 1  US-10-632-706-159  Sequence 159, Application US/10632706  PUDITION NO. US20040175385A1  GENERAL INFORMATION:  APPLICANT: MARKS, JAMES D.  ETHILB OF INVENTION: METHER D.  CURRENT APPLICATION NUMBER: US/10/632,706  CURRENT APPLICATION NUMBER: US 60/400,721  PRIOR APPLICATION NUMBER: US 60/400,721  PRIOR FILING DATE: 2002-08-01  PRIOR FILING DATE: 2002-08-31  INUMBER OF SEQ ID NOS: 278  SOFTWARE: PACTURE:  ORGANISM: Artificial  PEATURE:  OTHER INFORMATION: single chain antibody fragment  US-10-632-706-159 | 28 29 90.6 394 4 US-10-425-145-359544 Sequence 359544, 29 29 90.6 422 6 US-11-097-143-318 Sequence 318, App 30 29 90.6 425 4 US-10-207-655-33 Sequence 318, App 32 29 90.6 495 4 US-10-207-655-33 Sequence 251, App 33 29 90.6 511 4 US-10-437-963-143624 Sequence 251, App 35 29 90.6 513 4 US-10-437-963-161859 Sequence 251, App 36 29 90.6 544 US-10-437-963-161859 Sequence 143624, 37 29 90.6 965 4 US-10-437-963-161859 Sequence 44, App 39 29 90.6 970 4 US-10-371-877-44 Sequence 201115, 39 29 90.6 986 4 US-10-282-122A-42808 Sequence 201115, 39 29 90.6 986 4 US-10-282-122A-59736 Sequence 201115, 39 29 90.6 1259 4 US-10-282-122A-5669 Sequence 59736, App 39 90.6 1259 4 US-10-282-122A-5669 Sequence 59736, App 30 90.6 1259 4 US-10-282-122A-5669 Sequence 59736, App 30 90.6 2590 4 US-10-437-963-137624 Sequence 57669, App 30 90.6 2590 4 US-10-437-963-201113 Sequence 201113, 45 29 90.6 4737 4 US-10-437-963-201113 Sequence 201113, Sequence 2011115, App 30 90.6 4737 4 US-10-437-963-201113 Sequence 2011115, App 30 90.6 4737 4 US-10-437-963-201116 Sequence 2011115, App 30 90.6 4737 4 US-10-437-963-201116 Sequence 2011115, App 30 90.6 4737 4 US-10-437-963-201116 Sequence 2011115, App 30 90.6 4737 4 US-10-437-963-201116 Sequence 2011115, App 30 90.6 4737 4 US-10-437-963-201116 Sequence 201116, |
|-------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|------------------------------------------------------------------------------------------------------------------------------------------------------------------------|---------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|--------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
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TITLE OF INVENTION: THERAPEUTIC MONOCLONAL ANTIBODIES THAT NEUTRALIZE BOTULINUM;
TITLE OF INVENTION: NEUROTOXINS
FILE REFERENCE: 407T-895120US
CURRENT APPLICATION UNMBER: US/10/632,706
CURRENT FILING DATE: 2003-08-01
PRIOR APPLICATION NUMBER: US 60/400,721
PRIOR PRICING DATE: 2002-08-01
PRIOR PRILING DATE: 2002-08-01
PRIOR PRILING DATE: 1988-08-31
NUMBER OF SEQ ID MOSE: 278
SEQ ID MOSE: 278
SEQ ID NO. 167
LENGTH. 6
   ; OTHER INFORMATION: single chain antibody fragment US-10-632-706-167
   RESULT 3
US-10-632-706-167
   ; TYPE: PRT
, ORGANISM: Artificial
; PEATURE:
; OTHER INFORMATION: single chain antibody fragment
US-10-632-706-163
  US-10-632-706-171
   Ś
 Sequence 171, Application US/10632706
Publication No. US20040175385A1
GENERAL INFORMATION:
APPLICANT: MARKS, JAMES D.
APPLICANT: AMERSDORFER, PETER
   Sequence 167, Application US/10632706
Publication No. US20040175385A1
GENERAL INFORMATION:
APPLICANT: WARKS, JAMES D.
APPLICANT: AMERSDORFER, PETER
  Query Match
Best Local
  PRIOR FILING DATE: 1998-08-31
NUMBER OF SEQ ID NOS: 278
SOFTWARE: PatentIn version 3.2
SEQ ID NO 163
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Best Local
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SOFTWARE: PatentIn version 3.2
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ORGANISM: Artificial
FEATURE:
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100.0%; Pred. No. 1.7e+06;
  100.0%; Score 32; DB 4;
100.0%; Pred. No. 1.7e+06;
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   0; Mismatches
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   Length 5;
   Length 5;
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   <u>,</u>
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   Gaps
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   0
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APPLICANT: Hyseq, Inc
TITLE OF INVENTION: NOVEL NUCLEIC ACIDS AND POLYPEPTIDES
FILE REFERENCE: 790CIP3/US
CURRENT APPLICATION NUMBER: US/10/450,763
CURRENT FILING DATE: 2003-06-11
PRIOR APPLICATION NUMBER: PCT/US01/08631
PRIOR FILING DATE: 2001-03-30
PRIOR FILING DATE: 2001-03-31
PRIOR PELING DATE: 2000-03-31
PRIOR FILING DATE: 2000-03-31
PRIOR PELING DATE: 2000-03-31
PRIOR APPLICATION NUMBER: 09/540,167
PRIOR APPLICATION NUMBER: 09/649,167
PRIOR FILING DATE: 2000-08-23
  Ś
  밁
  á
   OTHER INFORMATION: Clone ID: PAT_MRT3847_130375C.1.pep US-10-424-599-176475
  밁
  US-10-450-763-50856
  RESULT 5
US-10-424-599-176475
   US-10-632-706-171
   APPLICANT: La Rosa Thomas J
APPLICANT: Kovalic David K
APPLICANT: Zhou Yihua
APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Therebf for Plant Improvement
FILE REFERENCE: 38-21(5323)B
FILE REFERENCE: 38-21(5323)B
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 176475
LENGTH: 73
   Sequence 50856, Application US/10450763 Publication No. US20050196754A1 GENERAL INFORMATION:
  Sequence 176475, Application US/10424599 Publication No. US20040031072A1 GENERAL INFORMATION:
  SEQ ID NO 171
LENGTH: 5
SEQ ID NO 50856
   Matches
   Matches
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Best Local Similarity
               NUMBER OF SEQ ID NOS: 60736
SOFTWARE: Custom
  NAME/KEY: unsure
LOCATION: (1)..(73)
OTHER INFORMATION: unsure at all Xaa locations
   OTHER INFORMATION: single chain antibody fragment
   ORGANISM: Artificial FEATURE:
  TYPE: PRT
   FEATURE:
   FEATURE:
   ORGANISM: Glycine max
  TYPE: PRI
  Local Similarity
  47 DYDMH 51
  1 DYDMH 5
  1 DYDMH 5
   5; Conservative
   Conservative
   100.0%; Score 32; Di
100.0%; Pred. No. 61
tive 0; Mismatches
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   0
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  Length 5;
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   Gaps
   Gaps
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; TYPE: PRT ; ORGANISM: Homo sapiens US-10-450-763-50856

Matches

26 DYDMH 30

1 DYDMH 5

Query Match Best Local Similarity

similarity 100.0%; Similarity 100.0%; Similarity 0.0%;

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Mismatches

Score 32; DB 5; Pred. No. 1.2e+02;

Length 144;

0

Gaps

0

US-10-779-461-18
; Sequence 18, Application US/10779461
; Publication No. US20040166544A1
; GENERAL INFORMATION:

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Query Match
Best Local Similarity
****ches 5; Conserve
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   ; FEATURE:
; OTHER INFORMATION: phage display generated human antibody
US-10-779-461-18
   US-10-369-493-23241
S
  APPLICANT: Cao, Yongwei
APPLICANT: Hinkle, Gregory J.
APPLICANT: Slater, Steven C.
APPLICANT: Goldman, Barry S.
APPLICANT: Chen, Xianfeng
  NUMBER OF SEQ ID NOS: 161
SOFTWARE: PatentIn version 3.2
SEQ ID NO 18
LENGTH: 253
   PRIOR APPLICATION NUMBER: US 60/360,039
PRIOR FILING DATE: 2002-02-21
NUMBER OF SEQ ID NOS: 47374
SEQ ID NO 23241
LENGTH: 319
  Sequence 23241, Application US/10369493 Publication No. US20030233675A1
  GENERAL INFORMATION:
  Matches
  Query Match
Best Local Similarity
  TYPE: PRT ORGANISM: Bacillus subtilis-10-369-493-23241
   APPLICANT: MORTON, Philip A
TITLE OF INVENTION: ANTIBODIES TO C-MET FOR THE TREATMENT OF CANCERS
FILE REFERENCE: 009807.
CURRENT APPLICATION NUMBER: US/10/779,461
CURRENT FILING DATE: 2004-02-13
PRIOR APPLICATION NUMBER: 60/447,073
PRIOR APPLICATION NUMBER: 60/447,073
PRIOR FILING DATE: 2003-02-13
   TITLE OF INVENTION: EXPRESSION OF MICROBIAL PROTEINS IN PLANTS FOR PRODUCTION OF TITLE OF INVENTION: PLANTS WITH IMPROVED PROPERTIES FILE REPERRINCE: 38-10(52052)B
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CURRENT FILING DATE: 2003-02-28
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ORGANISM: artificial
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    1 DYDMH 5
   100.0%; ilarity 100.0%; Conservative 0
                                      Conservative
                                      100.0%; Score 32; DB 4; Length 319; 100.0%; Pred. No. 2.5e+02; ive 0; Mismatches 0; Indels
  0
   Score 32; DB 4; Length 253; Pred. No. 2e+02;
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  0
  Indels
  0
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  Gaps
  Gaps
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RESULT 11 US-10-514-150-10

; Sequence 10,

Application US/10514150

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; ORGANISM: Pyrococcus horikoshii
US-10-369-493-20409
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   ; SEQ ID NO 1346
LENGTH: 465
; TYPE: PRT
; ORGANISM: Pyrococcus horikoshii
US-10-369-493-1346
   RESULT 10
US-10-369-493-1346
  밁
  Ś
                             á
뭉
  Sequence 1346, Application US/10369493 Publication No. US20030233675A1 GENERAL INFORMATION:
  Sequence 20409, Application US/10369493 Publication No. US20030233675A1
   SEQ ID NO 20409
  GENERAL INFORMATION:
  Query Match
Best Local Similarity
Matches 5; Conserv
   Matches
   Query Match
  APPLICANT: Cao, Yongwei
APPLICANT: Hinkle, Gregory J.
APPLICANT: Slater, Steven C.
APPLICANT: Slater, Steven C.
APPLICANT: Goldman, Barry S.
APPLICANT: Goldman, Barry S.
TITLE OF INVENTION: EXPRESSION OF MICROBIAL PROTEINS IN PLANTS FOR PRODUCTION OF TITLE OF INVENTION: PLANTS WITH IMPROVED PROPERTIES
FILE REFERENCE: 38-10(52052)B
CURRENT APPLICATION NUMBER: US/10/369,493
CURRENT APPLICATION NUMBER: US/10/369,493
  APPLICANT: Cao, Yongwei
APPLICANT: Hinkle, Gregory J.
APPLICANT: Slater, Steven C.
APPLICANT: Goldman, Barry S.
APPLICANT: Chen, Xianfeng
  CURRENT APPLICATION NUMBER: US/10/369,493
CURRENT FILING DATE: 2003-02-28
PRIOR APPLICATION NUMBER: US 60/360,039
PRIOR FILING DATE: 2002-02-21
NUMBER OF SEQ ID NOS: 47374
  PRIOR APPLICATION NUMBER: US 60/360,039 PRIOR FILING DATE: 2002-02-21
  NUMBER OF SEQ ID NOS: 47374
   TITLE OF INVENTION: EXPRESSION OF MICROBIAL PROTEINS IN PLANTS FOR PRODUCTION OF TITLE OF INVENTION: PLANTS WITH IMPROVED PROPERTIES FILE REFERENCE: 38-10(52052)B
   LENGTH: 457
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   235 DYDMH 239
241 DYDMH 245
   1 DYDMH 5
                                      1 DYDMH 5
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100.0%; Pred. No. 3.5e+02;
tive 0; Mismatches 0;
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   Length 465;
   Indels
  Indels
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   Gaps
  Gaps
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Publication No. US20050233957A1

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APPLICANT: John Nicholas WOOD
APPLICANT: Misbah MALIK-HALL
TITLE OF INVENTION: SODIUM CHANNEL REGULATORS AND MODULATORS
FILB REFERENCE: 117-528 / N.88745B GCW
CURRENT APPLICATION NUMBER: US/10/514/150
CURRENT FILING DATE: 2004-11-12
PRIOR APPLICATION NUMBER: PCT/GB03/02225
PRIOR APPLICATION NUMBER: PCT/GB03/02225
PRIOR APPLICATION NUMBER: GB 0211833.9
PRIOR FILING DATE: 2002-05-22
PRIOR FILING DATE: 2002-05-22
NUMBER OF SEQ ID NOS: 10
SOFTWARE: PATENTIN VETSION 3.1
SEQ ID NO 10
LENGTH: 564
TYPE: PRT
OPGARMISM. HOTO SERVICES
RESULT 13
US-09-925-302-801
j Sequence 801, Application US/09925302
j Publication No. US20030064072A9
j GENERAL INFORMATION:
  뭐
  ; ORGANISM: Homo sapiens
; PERTURE;
; NAME/KEY: SITE
; LOCATION: (1)
; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
US-09-925-302-801
   RESULT 12
US-09-925-302-801
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   US-10-514-150-10
   Query Match
Best Local Similarity lov.
"""ches 5; Conservative
   NUMBER OF SEQ ID NOS: 896
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 801
LENGTH: 581
TYPE: PRT
   GENERAL INFORMATION:
APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins and Antibodies
   Matches
  Query Match
Best Local Similarity
  Sequence 801, Application Patent No. US20020044941A1
   FILE REFERENCE: PA104
CURRENT APPLICATION NUMBER: US/09/925,302
CURRENT FILING DATE: 2001-08-10
PRIOR APPLICATION NUMBER: PCT/US00/05918
PRIOR PILING DATE: 2000-03-08
PRIOR APPLICATION NUMBER: 60/124,270
PRIOR PILING DATE: 1999-03-12
  ORGANISM: Homo sapiens
  PPLICANT: UNIVERSITY COLLEGE LONDON
  HWAYA 65
   22 DYDMH
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  1 DYDMH 5
   Mark BAKER
   Kenji
   Application US/09925302
   Conservative
   26
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100.0%; Pred. No. 4.3e+02;
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GENERAL INPORMATION:

APPLICANT: Hyseq, Inc

ITILE OF INVESTION: NOVEL NUCLEIC ACIDS AND POLYPEPTIDES

FILE REFERENCE: 790CIP3/US

CURRENT APPLICATION NUMBER: US/10/450,763

CURRENT FILING DATE: 2003-06-11

PRIOR APPLICATION NUMBER: PCT/US01/08631

PRIOR APPLICATION NUMBER: PCT/US01/08631

PRIOR PILING DATE: 2001-03-30

PRIOR PILING DATE: 2001-03-31

PRIOR APPLICATION NUMBER: 09/540,217

PRIOR APPLICATION NUMBER: 09/540,217

PRIOR PILING DATE: 2000-03-31

PRIOR APPLICATION NUMBER: 09/649,167

PRIOR FILING DATE: 2000-08-23

NUMBER OF SEQ ID NOS: 60736

SEPTWARE: CUSCOM
  ; FRATURE:
NAME/KBY: misc feature
; LOCATION: (1)...(585)
; OTHER INFORMATION: Xaa = X or * as defined in Table
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  Ś
  S
   밁
RESULT 15
  US-10-450-763-42152
   US-09-925-302-801
   APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins an
FILE REFERENCE: PA104
CURRENT APPLICATION NUMBER: US/09/925,302
CURRENT FILING DATE: 2001-08-10
PRIOR APPLICATION NUMBER: PCT/US00/05918
PRIOR FILING DATE: 2000-03-08
PRIOR APPLICATION NUMBER: 60/124,270
PRIOR APPLICATION NUMBER: 60/124,270
PRIOR FILING DATE: 1999-03-12
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  Matches
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Best Local (
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   Sequence 42152, Application US/10450763 Publication No. US20050196754A1
   Best Local Similarity
   Query Match
   NUMBER OF SEQ ID NOS: 896
SOFTWARE: Patentin Ver. 2.0
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NAME/FOR: SITE
LOCATION: (1)
OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
  LENGTH: 581
TYPE: PRT
ORGANISM: Homo sapiens
   ORGANISM: Homo sapiens
  ENGTH: 585
   Local Similarity
les 5; Conserv
   22 DYDMH
   39 DYDMH 43
  1 DYDMH 5
  1 DYDMH 5
  5
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   100.0%;
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Pred. No. 4.4e+02;
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US-10-450-763-34775, Application US/10450763; Sequence 34775, Application US/10450763; Publication No. US20050196754A1; GENERAL INFORMATION:

APPLICANT: Hyseq, Inc

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FILE OF INVENTION: NOVEL NUCLEIC ACIDS AND POLYPEPTIDES

FILE REFERENCE: 790C1P3/US

CURRENT APPLICATION NUMBER: US/10/450,763

CURRENT FILING DATE: 2003-66-11

PRIOR APPLICATION NUMBER: PCT/US01/08631

PRIOR APPLICATION NUMBER: 99/540,217

PRIOR APPLICATION NUMBER: 09/540,217

PRIOR APPLICATION NUMBER: 09/549,167

PRIOR APPLICATION NUMBER: 09/649,167

PRIOR FILING DATE: 2000-08-23

NUMBER OF SEQ ID NOS: 60736

SOCTWARE: CUSTOM

SEQ ID NO 34775

LENGTH: 596

TYPE: PRT

ORGANISM: Homo sapiens

FEATURE:

NAME/KEY: misc_feature

IOCATION: (1):..(596)

OTHER INFORMATION: Xaa = X or * as defined in Table 2

US-10-450-763-34775

Query Match

Best Local Similarity 100.0%; Score 32; DB 5; Length 596;

Best Local Similarity 100.0%; Pred. No. 4.5e+02;

Matches 5; Conservative 0; Mismatches 0; Indels 0; Gaps

Oy 1 DYDMH 5

Db 22 DYDMH 26

Search completed: November 21, 2005, 12:33:39

Job time: 8.19048 secs
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Result
No.
   Sequence:
   Title:
Perfect score:
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  Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
  Minimum DB seq length: 0
Maximum DB seq length: 200000000
   Scoring table:
   Run on:
  Database
   Total number of hits satisfying chosen parameters:
  protein -
 Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
   Score
 protein search, using sw model
  Published_Applications_AA_New:*

1: /cgn2_6/ptodata/2/pubpaa/USIO_NEW_PUB.pep:*

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  BLOSUM62
Gapop 10.0 , Gapext 0.5
   US-10-632-706-159
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   Match
 GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
   Length
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US-10-522-883-4

US-10-522-883-13

US-10-522-883-13

US-10-522-883-21

US-10-522-883-21

US-10-522-883-21

US-10-522-883-21

US-10-131-826A-336-100

US-10-131-826A-436

US-10-131-826A-436

US-10-131-826A-436

US-10-131-826A-436

US-11-179-977-8

US-11-179-977-8

US-11-174-248-48

US-11-174-248-48

US-11-174-248-48

US-11-174-248-48

US-11-144-248-48
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| RESULT 2  Sequence 19, Ap  Sequence 19, Ap  Publication No.  GENERAL INFORM  APPLICANT: CYT  TITLE OF INVEN  FILE REFERENCE  CURRENT APPLIC  CURRENT APPLIC  CURRENT SPLIC  CURRENT PILLNG  MUMBER OF 5E0  SOFTWARE: Pate  SEQ ID NO 19  LENGTH: 173  TYPE: PRT  ORGANISM: Art  FEATURE:  OTHER INFORMA | Query Matc<br>Best Local<br>Matches<br>Qy  | RESULT 1 US-10-522-883-2 Sequence 2, App. Publication No. GEMERAL INFORMA FILE OF INVEN FILE REFERENCE CURRENT FILING CURRENT FILING NUMBER OF SEQ NUMBER OF SEQ SOFTWARE: Pate SEQ ID NO 2 LENGTH: 153 TYPE: PRT ORGANISM: Art FEATURE: FEATURE: OTHER INFORMA US-10-522-883-2 |            | 22222222222222222222222222222222222222                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         |
|---------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|--------------------------------------------|---------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|------------|--------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| a a sa sa asa s                                                                                                                                                                                                                                                                                         | h<br>Simi<br>3;<br>1 DYD<br>1  : <br>5 DFD | BESSET TERMS                                                                                                                                                                                                                                                                    |            | 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 688 22 |
| a < 0 a                                                                                                                                                                                                                                                                                                 | 78. larity 60. Conservative MH 5 :  LH 79  | it.<br>31W<br>17 NU<br>18 ver                                                                                                                                                                                                                                                   |            |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                |
|                                                                                                                                                                                                                                                                                                         | 78.1%;<br>60.0%;<br>tive                   | 7 30 C PO                                                                                                                                                                                                                                                                       |            | 299<br>330<br>444<br>1944<br>11827<br>7<br>1857<br>7<br>1857<br>7<br>13<br>13<br>13<br>13<br>13<br>13<br>13<br>13<br>13<br>13<br>13<br>13<br>13                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                |
| 83<br>8ta<br>/52                                                                                                                                                                                                                                                                                        | score<br>pred.<br>2; Min                   | 0522883<br>1A1<br>ug substance,<br>US/10/522,88<br>02-02<br>3.1<br>3.1                                                                                                                                                                                                          | ALIC       | US-11-0<br>US-10-5<br>US-10-1<br>US-11-0<br>US-11-0<br>US-11-0<br>US-11-0<br>US-11-0<br>US-11-0<br>US-11-0<br>US-11-0<br>US-11-0<br>US-11-0<br>US-11-0<br>US-11-0<br>US-11-0<br>US-11-0<br>US-11-0<br>US-11-0                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  |
| Ψ                                                                                                                                                                                                                                                                                                       | re 25; DB<br>d. No. 6.1<br>Mismatches      | 88 e .                                                                                                                                                                                                                                                                          | ALIGNMENTS | 74-176- 10-386- 110-386- 31-826A 57-058- 57-058- 57-058- 57-058- 57-058- 57-058- 57-058- 57-058- 57-058- 57-107- 31-826A 31-826A 31-826A                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       |
| composition,                                                                                                                                                                                                                                                                                            | , 1,                                       | composition,                                                                                                                                                                                                                                                                    |            | \$\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        |
|                                                                                                                                                                                                                                                                                                         | Length 15<br>; Indels                      |                                                                                                                                                                                                                                                                                 |            |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                |
| preparation                                                                                                                                                                                                                                                                                             | 0,                                         | preparation                                                                                                                                                                                                                                                                     |            | Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      |
| 1 and us                                                                                                                                                                                                                                                                                                | Gарв                                       | and                                                                                                                                                                                                                                                                             |            | 198<br>498<br>664<br>662<br>661<br>20<br>20<br>20<br>20                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        |
| 60<br>80<br>•                                                                                                                                                                                                                                                                                           | 0;                                         | uses.                                                                                                                                                                                                                                                                           |            | App<br>App<br>App<br>App<br>App<br>Appl<br>Appl<br>Appl<br>Appl                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                |

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Ś
US-10-522-883-13
, Sequence 13, Application US/10522883
, Publication No. US20050249701A1
, GENERAL INFORMATION:
   ; FEATURE:
; OTHER INFORMATION: r-hIL-7 cDNA
US-10-522-883-4
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  밁
   용
  S
  US-10-522-883-4
   US-10-522-883-23
   RESULT 5
   US-10-522-883-23
  Sequence 23, Application US/10522883

Publication No. US20050249701A1

GENERAL INFORMATION:
APPLICANT: CYTHERIS

TITLE OF INVENTION: IL-7 drug substance, composition, preparation and uses.
FILE REFERENCE: B0131WO
CURRENT APPLICATION NUMBER: US/10/522,883

CURRENT PILING DATE: 2005-02-02

NUMBER OF SEQ ID NOS: 23

SOFTWARE: Patentin version 3.1

SEQ ID NO 23

LENGTH: 173
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CURRENT FILING DATE: 2005-02-02
NUMBER OF SEQ ID NOS: 23
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Publication No. US20050249701A1
GENERAL INFORMATION:
APPLICANT: CYTHERIS
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  TYPE: PRT
ORGANISM: Artificial sequence
   ORGANISM: Artificial sequence FEATURE:
   TYPE: PRT
   LENGTH: 177
  OTHER INFORMATION: HMM38PS-sIL-7
  Local Similarity
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  95 DFDLH 99
  95 DFDLH 99
   99 DFDLH 103
   1 DYDMH 5
  1 DYDMH 5
  1 DYDMH 5

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   Conservative
   Conservative
   78.1%;
  78.1%;
   78.1%;
   2
  2
   N
-
  Score 25;
Pred. No.
   Score 25;
Pred. No.
   Score 25; DB 1;
Pred. No. 7.1;
  Mismatches
  Mismatches
   Mismatches
   멂
  DB 1;
   ۲.
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  Length 177;
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US-10-522-883-21
Sequence 21, Application US/10522883
Publication No. US20050249701A1
GENERAL INFORMATION:
APPLICANT: CYTHERIS
TITLE OF INVESTION: IL-7 drug subst.
FILE REFERENCE: B0131WO
   밁
  ঠ
  밁
  Ś
   RESULT 6
US-10-522-883-17
  US-10-522-883-13
; OTHER INFORMATION: EPOPS-8IL-7 cDNA US-10-522-883-21
   RESULT 7
   OTHER INFORMATION: EPOPS-r-hIL-7 cDNA US-10-522-883-17
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Matches 3
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Publication No. US20050249701A1
GENERAL INFORMATION:
APPLICANT: CYTHERIS
   CURRENT APPLICATION NUMBER: US/10/522,883
CURRENT FILING DATE: 2005-02-02
NUMBER OF SEQ ID NOS: 23
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TYPE: PRT
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CURRENT FILING DATE: 205-02-02
NUMBER OF SEQ ID NOS: 23
SOFTWARE: PatentIn version 3.1
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LENGTH: 179
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LENGTH: 179
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   TITLE OF INVENTION: IL-7 drug substance, composition, preparation and uses. FILE REFERENCE: B0131WO
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                            TYPE: PRT
ORGANISM: Artificial sequence
FEATURE:
  TYPE: PRT ORGANISM: Artificial sequence
   OTHER INFORMATION: PS-sIL-7 cDNA
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  FEATURE:
  Local Similarity
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   99 DFDLH 103
   1 DYDMH 5
  1 DYDMH 5
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3; Conserv
   Conservative
   Conservative
   IL-7 drug substance, composition, preparation and uses.
  IL-7 drug substance, composition, preparation
  78.1%;
   78.1%; Score 25; E 60.0%; Pred. No. 7.
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  Score 25; DB
Pred. No. 7.1;
  Mismatches
   Mismatches
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  DB 1;
  0;
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  Length 177;
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  Indels
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   0
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101 DFDLH 105

1 DYDMH 5

Query Match Best Local Similarity Watches 3; Conservi

Conservative

2; Mismatches Score 25; DB 1; Pred. No. 7.2;

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Gaps

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Length 179;

78.1%; 60.0%;

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US-10-131-826A-332
; Sequence 332, Application US/10131826A
; Publication No. US20050245730A1
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  Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 550
SEQ ID NO 332
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  APPLICANT:
  TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P3330R1C128 CURRENT APPLICATION NUMBER: US/10/131,826A CURRENT FILING DATE: 2002-04-24 PRIOR APPLICATION NUMBER: 60/049911 PRIOR PELLING DATE: 1997-06-18 PRIOR PILING DATE: 1997-06-18 PRIOR APPLICATION NUMBER: 60/056974 PRIOR APPLICATION NUMBER: 60/056974 PRIOR APPLICATION NUMBER: 60/056974 PRIOR APPLICATION NUMBER: 60/056974 PRIOR APPLICATION NUMBER: 60/056974 PRIOR APPLICATION NUMBER: 60/056974 PRIOR APPLICATION NUMBER: 60/056974 PRIOR APPLICATION NUMBER: 60/056974
  PRIOR APPLICATION NUMBER: 60/059588
PRIOR FILING DATE: 1997-09-19
  APPLICANT: Baker,Kevin P.
APPLICANT: Beresini,Maur
  APPLICANT:
   PRIOR APPLICATION NUMBER: 60/059352
PRIOR FILING DATE: 1997-09-19
   PRIOR APPLICATION NUMBER: 60/
PRIOR FILING DATE: 1997-09-18
   APPLICANT:
   PRIOR APPLICATION NUMBER: 60/059122 PRIOR FILING DATE: 1997-09-17
  PRIOR APPLICATION NUMBER: 60
PRIOR FILING DATE: 1997-09-1
PRIOR APPLICATION NUMBER: 60
   APPLICANT:
  APPLICANT
  RIOR APPLICATION NUMBER: 60/059184
  RIOR FILING DATE: 1997-09-17
   RIOR FILING DATE: 1997-09-17
   LENGTH: 552
TYPE: PRT
   Local Similarity es 3; Conserv
  FILING DATE: 1997-09-17
   INFORMATION:
268 DYELH 272
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   Stewart, Timothy A. Tumas, Daniel . Watanabe, Colin K
  Goddard, Audrey
Godowski, Paul J.
Gurney, Austin L.
  Gao, Wei-Qiang
   Smith, Victoria
  Pilvaroff, Ellen
  Zhang, Zemin
  Gerritsen, Mary E.
  DeForge, Laura
   Wood,William
   Sherwood, Steven
  Beresini,Maureen
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60.0%;
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   60/059263
  60/059117
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; ORGANISM: Corynebacterium glutamicum
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PRIOR PELLING DATE: 2000-06-23
PRIOR PELLING DATE: 1999-06-25
PRIOR PELLING DATE: 1999-06-25
PRIOR APPLICATION NUMBER: US 60/143262
PRIOR PELLING DATE: 1999-07-09
PRIOR APPLICATION NUMBER: US 60/151281
PRIOR APPLICATION NUMBER: US 60/151281
PRIOR PILLING DATE: 1999-08-27
PRIOR APPLICATION NUMBER: DE 19930487.4
PRIOR APPLICATION NUMBER: DE 19930489.0
PRIOR APPLICATION NUMBER: DE 19930489.0
PRIOR FILLING DATE: 1999-07-01
PRIOR PILLING DATE: 1999-07-01
PRIOR PILLING DATE: 1999-07-01
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  Remaining Prior Application data removed - See File Wrapper or PALM. NUMBER OF SEQ ID NOS: 446
SEQ ID NO 290
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   Sequence 290, A Publication No.
  Sequence 100, Application US/10510386 Publication No. US20050244922A1
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  SOFTWARE: PatentIn version 3.3 SEQ ID NO 100
  GENERAL INFORMATION:
   APPLICANT: Jorgensen, Steen Troels
APPLICANT: Olsen, Peter Bjarke
APPLICANT: Rasmussen, Michael Dolberg
TITLE OF INVENTION: Improved Bacillus Host Cell
FILE REFERENCE: 10294_204_US
CURRENT APPLICATION NUMBER: US/10/510,386
CURRENT FILING DATE: 2004-10-04
  APPLICANT: Zelder, Oskar
APPLICANT: Haberhauer, Gregor
TITLE OF INVENTION: CORYNEBACTERIUM GLUTAMICUM GENES ENCODING PROTEINS
TITLE OF INVENTION: INVOLVED IN MEMBRANE SYNTHESIS AND MEMBRANE
TITLE OF INVENTION: TRANSPORT
FILE REFERENCE: BGI-131CPCN
FILE REFERENCE: BGI-131CPCN
  APPLICANT: Pompejus, Markus APPLICANT: Kroger, Burkhare
   CURRENT APPLICATION NUMBER: US/11/082,389
CURRENT FILING DATE: 2005-03-16
   NUMBER OF SEQ ID NOS: 248
  APPLICANT: Andersen, Jens Tonne
APPLICANT: Clausen, Ib Groth
APPLICANT: Jorgensen, Steen Tro
APPLICANT: Olsen, Peter Bjarke
   PRIOR APPLICATION NUMBER: DE 19941379.7 PRIOR FILING DATE: 1999-08-31
                      LENGTH: 27
TYPE: PRT
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  APPLICATION NUMBER: DE 19931549.3 FILING DATE: 1999-07-08 APPLICATION NUMBER: DE 19931550.7 FILING DATE: 1999-07-08
   FILING DATE: 1999-07-09
  APPLICATION NUMBER: DE 19932134.5
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  Kroger, Burkhard
  Zelder, Oskar
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100.0%; Pred. No.
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  US-10-510-386-100
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Best Local Similarity 50.0
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NUMBER OF SEQ ID NOS: 550
SEQ ID NO 136
LENGTH: 278
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APPLICANT:
APPLICANT:
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CURRENT FILING DATE: 2002-04-24
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PRIOR FILING DATE: 1997-06-18
  PRIOR APPLICATION NUMBER: 60/056974
PRIOR FILING DATE: 1997-08-26
PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
  APPLICANT: Zhang, Zemin
TITLE OP INVENTION: SECRETED AND TRANSMEMBRANB POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3330R1C128
  PRIOR FILING DATE: 1997-09-19
PRIOR APPLICATION NUMBER: 60/059588
PRIOR FILING DATE: 1997-09-19
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   APPLICANT:
   APPLICANT: Baker, Kevin
  PPLICANT:
   PPLICANT:
  PPLICANT:
   Local Similarity
  APPLICATION NUMBER: 60/059115
FILING DATE: 1997-09-17
APPLICATION NUMBER: 60/059117
  APPLICATION NUMBER: 60/059184
  APPLICATION NUMBER: 60/059352
   APPLICATION NUMBER: 60/059263
  FILING DATE: 1997-09-17
   APPLICATION NUMBER: 60/059122
  FILING DATE: 1997-09-17
   FILING DATE: 1997-09-18
109 DYSLH 113
  86 DYDLY 90
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   1 DYDMH 5
   Goddard, Audrey
Godowski, Paul J.
Gurney, Austin L.
Sherwood, Steven
Smith, Victoria
  Watanabe, Colin K
Wood, William
  Filvaroff, Ellen
Gao, Wei-Qiang
  Tumas, Daniel
  Stewart, Timothy A.
   Gerritsen, Mary E.
   Beresini, Maureen
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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME

PILE REPERENCE: P3330R1C128

CURRENT APPLICATION UNMBER: US/10/131,826A

CURRENT FILING DATE: 2002-04-24

PRIOR APPLICATION NUMBER: 60/049911

PRIOR FILING DATE: 1997-06-18

PRIOR APPLICATION NUMBER: 60/05974

PRIOR APPLICATION NUMBER: 60/059113

PRIOR APPLICATION NUMBER: 60/059113

PRIOR APPLICATION NUMBER: 60/059113

PRIOR APPLICATION NUMBER: 60/059115

PRIOR APPLICATION NUMBER: 60/059117

PRIOR APPLICATION NUMBER: 60/059117

PRIOR APPLICATION NUMBER: 60/059117

PRIOR APPLICATION NUMBER: 60/059117

PRIOR FILING DATE: 1997-09-17

PRIOR FILING DATE: 1997-09-17

PRIOR APPLICATION NUMBER: 60/059117

PRIOR APPLICATION NUMBER: 60/059117

PRIOR APPLICATION NUMBER: 60/059117

PRIOR APPLICATION NUMBER: 60/059117

PRIOR APPLICATION NUMBER: 60/059112
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US-11-179-977-8
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US-10-131-826A-456
; Sequence 456, Application US/10131826A
; Publication No. US20050245730A1
  ; ORGANISM: Homo Sapien 
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SEQ ID NO 456
LENGTH: 293
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APPLICANT:
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   APPLICANT:
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  APPLICANT:
   APPLICATION NUMBER: 60/059263
FILING DATE: 1997-09-18
APPLICATION NUMBER: 60/059352
   FILING DATE: 1997-09-17
APPLICATION NUMBER: 60/059184
FILING DATE: 1997-09-17
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Wood, William
   Godowski, Paul J. Gurney, Austin L.
   Smith, Victoria
  Goddard, Audrey
  Gerritsen, Mary E
   Stewart, Timothy A.
   Sherwood, Steven
  Gao, Wei-Qiang
   DeForge, Laura
   Beresini, Maureen
  Filvaroff, Ellen
   Conservative
   71.9%;
   Score 23; bB 1;
Pred. No. 33;
   0; Mismatches
  Length 293;
   0,
   Gaps
   0
```

Sequence 8, Application US/11179977

Publication No. US20050249789A1

GENERAL INFORMATION:
APPLICANT: Genencor International, Inc.
APPLICANT: Genencor International, Inc.

```
US-10-632-150-26

| Sequence 26, Application US/10632150
| Publication No. US20050251871A1
| Publication No. US20050251871A1
| GENERAL INFORMATION:
| APPLICANT: Chiaux, D.
| APPLICANT: Latres, E.
| APPLICANT: Latres, E.
| TITLE OF INVENTION: NOVEL UPIQUITIN LIGASES AS THERAPEUTIC TARGETS
| FILE REFERENCE: 5914-081
| CURRENT APPLICATION NUMBER: US/10/632,150
| CURRENT FILING DATE: 2003-07-30
| PRIOR APPLICATION NUMBER: US/09/385,219
| PRIOR PILING DATE: 1999-08-27
| PRIOR PILING DATE: 1999-08-28
| PRIOR APPLICATION NUMBER: 60/018,568
| PRIOR APPLICATION NUMBER: 60/118,568
   ; FILE REFERENCE: GC511-PCT
; CURRENT APPLICATION NUMBER: US/11/179,977
; CURRENT FILING DATE: 2005-07-12
; NUMBER OF SEQ ID NOS: 21
; SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 8
; SEQ ID NO 8
; LENGTH: 296
; TYPE: PRT
; ORGANISM: Bacillus
US-11-179-977-8
   밁
  US-11-074-176-244
  NUMBER OF SEQ ID NOS: 381
SOFTWARE: FRSTSEQ for Windows Version 4.0
SEQ ID NO 244
LENGTH: 628
TYPE: PRT
  Sequence 244, Application US/11074176
Publication No. US20050250135A1
GENERAL INFORMATION:
   Query Match 71.9%; Score 23; DB 7; Length 628; Best Local Similarity 60.0%; Pred. No. 74; Matches 3; Conservative 2; Mismatches 0; Indels
   Matches
  Query Match
Best Local Similarity
   ORGANISM: Lactobacillus acidophilus -11-074-176-244
  FILE REFERENCE: 5051-694
CURRENT APPLICATION NUMBER: US/11/074,176
CURRENT FILING DATE: 2005-03-07
PRIOR APPLICATION NUMBER: 60/551,161
PRIOR FILING DATE: 2004-03-08
   APPLICANT: Altermann, Eric
APPLICANT: McAuliffe, Olivia
APPLICANT: Peril, Andrea Azcarate
TITLE OF INVENTION: Nucleic Acid Sequences Encoding
TITLE OF INVENTION: Stress-Related Proteins and Uses Therefore
  APPLICANT: Klaenhammer, Todd R. APPLICANT: Russell, William M. APPLICANT: Altermann, Eric APPLICANT: MCAuliffe, Olivia
APPLICATION NUMBER: 60/124,449
   190 YDLH 193
   |:|:|
145 DHDIH 149
   2 YDMH 5
   1 DYDMH 5

    Conservative

  71.9%;
75.0%;
  Score 23; DB : Pred. No. 34; 1; Mismatches
   7; Length 296;
   Indels
   <u>,</u>
  <u>.</u>
   Gaps
  Gaps
   0
  0
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; PRIOR FILING DATE: 1999-03-15
; NUMBER OF SEQ ID NOS: 90
; SOFTWARE: Patentin Ver. 2.0
; SEQ ID NO 26
; LENGTH: 634
; TYPE: PRT ...
; ORGANISM: Homo sapiens
; PEATURE:
; NAME/KEY: SITE
; LOCATION: all Xaa positions
; OTHER INFORMATION: Xaa=unknown amino acid residue
US-10-632-150-26

Query Match
Best Local Similarity 75.0%; Pred. No. 75;
Best Local Similarity 75.0%; Pred. No. 75;
Matches 3; Conservative 1; Mismatches 0; Gaps
Qy 2 YDMH 5
Qy 2 YDMH 5
Qh 621 YDLH 624
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Search completed: November 21, 2005, 12:33:52 Job time: 0.191575 secs

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Run
   Minimum DB seq length: Maximum DB seq length:
  Scoring table:
   Sequence:
   Perfect score:
  OM protein -
   Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
   Total number of hits satisfying chosen parameters:
  95:
  protein search, using sw model
   A_Geneseq_21:*
1: geneseqp1980
   BLOSUM62
   1 RFTISRDNSKNTLFLQMNSLRADDTAVYYCAR 32
  November 21,
  4.
   2443163 seqs, 439378781 residues
   Gapop 10.0 , Gapext 0.5
  US-10-632-706-196
   GenCore version 5.1.6 Copyright (c) 1993 - 2005 Compugen Ltd.
              geneseqp1980s:*
geneseqp2000s:*
geneseqp20001s:*
geneseqp2001s:*
geneseqp2003as:*
geneseqp2003as:*
geneseqp2003hs:*
geneseqp2005s:*
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  2005, 11:49:32 ;
  ; Search time 42.8425 Seconds (without alignments) 328.182 Million cell updates/sec
  2443163
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

| Result<br>No.<br>1<br>2<br>3<br>3<br>4<br>4<br>7<br>7 | Score<br>165<br>165<br>165<br>165<br>162<br>162 | Query<br>Match<br>100.0<br>100.0<br>100.0<br>98.2<br>98.2<br>98.2<br>98.2 | Length 32 32 32 32 31 31 116 117 | . 8 8 8 8 4 6 5 6 | ID ADR38791 ADR38794 ADR388797 ADR38800 AAU02529 AAU02529 AABG30486 ABG30486 ABG30486 ABG30486 | Description<br>Adr38791 Mo<br>Adr38794 Mo<br>Adr388797 Mo<br>Adr38800 Mo<br>Adr38806 Mo<br>Adr02529 An<br>Abg30486 Hu<br>Abg30486 Hu<br>Abg30486 Hu |
|-------------------------------------------------------|-------------------------------------------------|---------------------------------------------------------------------------|----------------------------------|-------------------|------------------------------------------------------------------------------------------------|-----------------------------------------------------------------------------------------------------------------------------------------------------|
| ω N                                                   | 165<br>165                                      |                                                                           | 32<br>32                         | <b>co</b> co      | ADR38794<br>ADR38797                                                                           |                                                                                                                                                     |
| .4.                                                   | 165                                             |                                                                           | 32                               | œ                 | ADR38800                                                                                       |                                                                                                                                                     |
| ഗ                                                     | 162                                             |                                                                           | 113                              | 4.                | AAU02529                                                                                       |                                                                                                                                                     |
| o.                                                    | 162                                             |                                                                           | 116                              | S                 | ABG30486                                                                                       |                                                                                                                                                     |
| 7                                                     | 162                                             |                                                                           | 117                              | ហ                 | ABG30484                                                                                       |                                                                                                                                                     |
| œ                                                     | 162                                             |                                                                           | 118                              | 9                 | ADW08865                                                                                       |                                                                                                                                                     |
| 9                                                     | 162                                             |                                                                           | 126                              | 4                 | AAG93640                                                                                       |                                                                                                                                                     |
| 10                                                    | 162                                             |                                                                           | 126                              | σ                 | ABO27447                                                                                       |                                                                                                                                                     |
| 11                                                    | 162                                             | 98.2                                                                      | 127                              | 4                 | AAG93627                                                                                       |                                                                                                                                                     |
| 12                                                    | 162                                             | 98.2                                                                      | 127                              | 0                 | ABO27434                                                                                       |                                                                                                                                                     |
| 13                                                    | 162                                             | 98.2                                                                      | 244                              | 7                 | ADG30422                                                                                       |                                                                                                                                                     |
| 14                                                    | 161                                             |                                                                           | 115                              | œ                 | ADP22238                                                                                       |                                                                                                                                                     |
| 15                                                    | 161                                             |                                                                           | 121                              | 8                 | AD036356                                                                                       |                                                                                                                                                     |
| 16                                                    | 161                                             |                                                                           | 121                              | æ                 | ADQ75229                                                                                       |                                                                                                                                                     |
| 17                                                    | 161                                             |                                                                           | 122                              | œ                 | ADP22292                                                                                       |                                                                                                                                                     |
| 18                                                    | 161                                             |                                                                           | 122                              | œ                 | ADP22308                                                                                       |                                                                                                                                                     |
| 19                                                    | 161                                             |                                                                           | 123                              | 7                 | ADL91329                                                                                       |                                                                                                                                                     |
| 20                                                    | 161                                             | 97.6                                                                      | 463                              | ω                 | AAY93701                                                                                       |                                                                                                                                                     |
| 21                                                    | 161                                             |                                                                           | 463                              | ω                 | AAY93727                                                                                       |                                                                                                                                                     |
| 22                                                    | 161                                             |                                                                           | 463                              | w                 | AAY93728                                                                                       |                                                                                                                                                     |
| 23                                                    | 161                                             | 97.6                                                                      | 463                              | σ                 | AAE35882                                                                                       |                                                                                                                                                     |
| •                                                     | 7                                               | )                                                                         |                                  |                   |                                                                                                |                                                                                                                                                     |

The invention describes an isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a specific clone where (I) binds to and neutralises botulinum neurotoxin type A (BoNT/A). An isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a clone chosen from clone 25, C25, C39, IC6, 3D12, B4, IF3, huC25, A1, Ar2, WR1(V), WR1(T), 3-1, 3-8, 3-10 and ING1, where (I) binds to and neutralizes botulinum neurotoxin type A (BoNT/A). Also described are: a polypeptide (II) comprising BoNT/A neutralising epitope having an epitope that is

Example 4; SEQ ID NO 193; 110pp; English.

New isolated antibody that neutralizes botulinum neurotoxin type A, useful for diagnosing botulism or for treating pathologies associated with botulinum neurotoxin poisoning.

WPI; 2004-652009/63.

Marks JD,

Amersdorfer P;

(REGC ) UNIV CALIFORNIA.

| 4.5       | 44       | 43       | 42        | 41       | 40        | 39       | 38       | 37       | 36        | 35       | 34       | 33       | 32        | 31        | 30        | 29       | 28       | 27       | 26       | 25        |
|-----------|----------|----------|-----------|----------|-----------|----------|----------|----------|-----------|----------|----------|----------|-----------|-----------|-----------|----------|----------|----------|----------|-----------|
| 158       | 158      | 158      | 158       | 158      | 158       | 158      | 158      | 158      | 158       | 158      | 158      | 158      | 158       | 158       | 158       | 158      | 158      | 158      | 158      | 159       |
|           |          | 95.8     |           |          |           |          |          |          |           |          |          |          |           |           |           |          |          |          | 95.8     |           |
| 32        | 32       | 32       | 32        | 32       | 32        | 32       | 32       | 32       | 32        | 32       | 32       | 32       | 32        | 32        | 32        | 32       | 32       | 32       | 32       | 120       |
| 9         | ø        | 9        | 9         | 9        | 9         | 9        | 9        | 9        | 9         | 9        | 9        | ဖ        | 9         | æ         | æ         | œ        | ψ        | U        | N        | 4         |
| ABA21479  | ADY31530 | ADY31374 | ADY31392  | ADY31506 | ADY31371  | ADY31398 | ADY31503 | ADY31524 | ADX27039  | ADW80297 | ADW80310 | ADW96738 | ADW96725  | ADQ90726  | ADL93671  | ADL93533 | AAU70472 | AAB19693 | AAR87051 | AAU02508  |
| Aea21479  | Ady31530 | Ady31374 | Ady31392  | Ady31506 | Ady31371  | Ady31398 | Ady31503 | Ady31524 | Adx27039  | Adw80297 | Adw80310 | AGW96738 | Adw96725  | Adq90726  | Ad193671  | Ad193533 | Aau70472 | Aae19693 | Aar87051 | Aau02508  |
| Human and | _        | _        | Human ant | _        | Human ant | _        | _        | -        | Human gro |          |          |          | Anti-EGFR | Anti-VEGE | Human CD4 |          |          |          |          | Anti-adip |

## ALIGNMENTS

## ADR3871 ADR3873 XX ADR3873 XX ADR3873 XX ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 ADR37 A 31-AUG-1998; 98US-00144886. 01-AUG-2002; 2002US-0400721P. 01-AUG-2003; 2003US-00632706. US2004175385-A1. Mus sp. antibacterial; antibody; botulinum neurotoxin type A; BoNT/A; BoNT/A neutralising epitope; anti-botulinum neurotoxin antibody; toxin neutralisation; botulinum neurotoxin poisoning; mouse; heavy chain; Mouse heavy chain anti-BoNT-antibody framework 3 seqid 193. 02-DEC-2004 ADR38791 standard; peptide; 32 AA. 09-SEP-2004. ADR38791; (first entry)

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RESULT 2
ADR38794
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  specifically bound by an antibody expressed by clones as mentioned in (I) producing (I); and a composition (III) comprising several anticodition producing (I); and a composition (III) comprising several anticodition of a botulinum neurotoxin and the combination of cantibodies shows greater toxin neutralisation than the single antibodies in surplus. The following are disclosed; a pharmaceutical composition (C); and a kit comprising (I); and a kit comprising (I) is useful for neutralising comprising (I); and a kit comprising a botulinum neurotoxin which involves contacting neurotoxin with (I) in surplus, where each of (I) is specific for a different epitope of the botulinum neurotoxin and the combination of antibodies shows greater toxin neutralisation than the single antibodies in surplus. (I) is useful for diagnosing the botulism or for treating pathologies associated with botulinum neurotoxin poisoning. (I) exhibits specificity and affinity towards BONTA. (I) canables rapid detection or diagnosis of botulism. This is the amino acid of course of mouse heavy chain anti-BoNT-antibody framework region 3.
   Query Match
Best Local S
Matches 32
   Mus
  New isolated antibody that neutralizes botulinum neurotoxin type useful for diagnosing botulism or for treating pathologies associ
   31-AUG-1998; 98US-00144886.
01-AUG-2002; 2002US-0400721P.
   antibacterial; antibody; botulinum neurotoxin type A; BONT/A; BONT/A neutralising epitope; anti-botulinum neurotoxin antibody; toxin neutralisation; botulinum neurotoxin poisoning; mouse; heavy chain;
   Sequence 32 AA;
   WPI;
   01-AUG-2003; 2003US-00632706.
   09-SEP-2004.
   US2004175385-A1
  Mouse heavy chain anti-BoNT-antibody framework 3 seqid 196.
  02-DEC-2004 (first entry)
   ADR38794 standard; peptide;
   (REGC ) UNIV CALIFORNIA.
   2004-652009/63.
  à
  32;
  -
   1 RETISRONSKNTLFLOMNSLRADDTAVYYCAR 32
  Similarity
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   Amersdorfer
   Conservative
  98US-00144886.
  100.0%;
   32
   .
  Score
Pred.
   Mismatches
  165; DB 8;
No. 9.9e-16;
  32
  Length
  Indels
   32
   0
   Gaps
   0
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The invention describes an isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a specific clone where (I) binds to and neutralises botulinum neurocoxin type A (BONT/A). An isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a clone chosen from clone specifically bound by an antibody expressed by a clone chosen from clone S25, C25, C39, 1C6, 3D12, B4, 1F3, huC25, Ar1, Ar2, WR1(V), WR1(T), 3-1, 3-8, 3-10 and ING1, where (I) binds to and neutralizes botulinum

The invention describes an isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a specific clone where (I) binds to and neutralises botulinum neurotoxin type A (BoNT/A). An isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a clone chosen from clone

Example

4; SEQ

ID NO 199; 110pp; English

Example 4;

SEQ ID NO 196; 110pp; English.

botulinum neurotoxin poisoning.

pathologies associated

```
ARRSULT 3
ADR38791
ID ADR33
XX ADR33
XX ADR33
XX ADR33
XX ADR33
XX BONT
DT 02-D
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DT 02-D
DT 02-D
DT 02-D
DT 02-D
DT 02-D
DT 02-D
DT 02-D
DT 01-A
Exam
DE MAIN
XX BONT
XX BONT
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  cc comprising BONT/A neutralising epitope having an epitope that is specifically bound by an antibody expressed by clones as mentioned in (I) producing (I); and a composition (III) comprising several anti-
cc producing (I); and a composition (III) comprising several anti-
cc botulinum neurotoxin antibodies, where each antibody is specific for a different epitope of a botulinum neurotoxin and the combination of comprising (I); and a kit comprising (I) (I) is useful for neutralisation than the single antibodies shows greater toxin neutralisation than the single antibodies comprising (I); and a kit comprising (I). (I) is useful for neutralising comprising (I) and a kit comprising (I) a botulinum neurotoxin which comprising (I) antibody and for neutralising a botulinum neutrotoxin and the combination of antibodies shows greater toxin neutralisation than the compliantion of antibodies shows greater toxin neutralisation than the compliantion of antibodies shows greater toxin neutralisation than the compliantion of antibodies shows greater toxin neutralisation than the compliantion of antibodies shows greater toxin neutralisation than the compliantion of antibodies shows greater toxin neutralisation than the compliantion of antibodies shows greater toxin neutralisation than the compliantion of antibodies shows greater toxin neutralisation than the compliantion of antibodies shows greater toxin neutralisation than the compliantion of antibodies shows greater toxin meutralisation than the compliantion of antibodies shows greater toxin meutralisation than the compliantion of antibodies shows greater toxin meutralisation than the compliantion of antibodies shows greater toxin meutralisation than the compliantion of antibodies shows greater toxin meutralisation than the compliantion of antibodies shows greater toxin meutralisation than the compliantion of antibodies shows greater toxin meutralisation than the compliantion of antibodies shows greater toxin meutralisation of the botulism of the compliantion of antibodies sho
   Query Match
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Matches 32
   New isolated antibody that neutralizes botulinum neurotoxin type A, useful for diagnosing botulism or for treating pathologies associated with botulinum neurotoxin poisoning.
  31-AUG-1998; 98US-00144886.
01-AUG-2002; 2002US-0400721P.
  antibacterial; antibody; botulinum neurotoxin type A; BONT/A;
BONT/A neutralising epitope; anti-botulihum neurotoxin antibody;
toxin neutralisation; botulinum neurotoxin poisoning; mouse; hea
   Sequence 32 AA;
  Marks
   01-AUG-2003; 2003US-00632706.
   09-SEP-2004.
   US2004175385-A1
   Mus sp
  Mouse heavy chain anti-BoNT-antibody framework 3 seqid 199.
  02-DEC-2004
  ADR38797;
  ADR38797 standard;
  (REGC ) UNIV CALIFORNIA.
  2004-652009/63
   32;
  1 RFTISRDNSKNTLFLQMNSLRADDTAVYYCAR 32
   h 100.0%;
Similarity 100.0%;
32; Conservative 0
   ω
  Amersdorfer P;
  (first entry)
  peptide;
  32
   0
   Score 165; DB 8;
Pred. No. 9.9e-16;
Mismatches 0;
  B
  neurotoxin poisoning; mouse; heavy chain;
   Length
   0
   Gaps
   0
```

```
CC 325, C25, C39, 1C6, 3D12, B4, 1F3, huC25, Ar1, Ar2, WR1(Y), WR1(T), 3-1, CC 3-8, 3-10 and ING1, where (I) binds to and neutralizes botulinum CC neurotoxin type A (BoNT/A). Also described are: a polypeptide (II) CC comprising BoNT/A neutralising epitope having an epitope that is CC specifically bound by an antibody expressed by clones as mentioned in (I) producing (I); and a composition (II) comprising several anticoccomposition (II) comprising several anticoccomposition (II) comprising several anticoccomposition (I) antibodies shows greater toxin neutralisation than the single antibodies in surplus. The following are disclosed: a pharmaceutical composition (CC comprising (I); and a kit comprising (I). (I) is useful for neutralising CC specific for a different epitope of the botulinum neurotoxin which (I) is specific for a different epitope of the botulinum neurotoxin and the CC combination of antibodies shows greater toxin neutralisation than the CC single antibodies in surplus. (I) is useful for diagnosing the botulism CC or for treating pathologies associated with botulinum neurotoxin composition (I) exhibits specificity and affinity towards BONT/A. (I) cenables rapid detection or diagnosis of botulism. This is the amino acid sequence of mouse heavy chain anti-BONT-antibody framework region 3.
  S
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  Matches
  Query Match
  BONT/A neutralising epitope; anti-botulinum neurotoxin antibody; toxin neutralisation; botulinum neurotoxin poisoning; mouse; heavy chain;
   Mouse heavy chain anti-BoNT-antibody framework 3 seqid 202
   02-DEC-2004 (first entry)
  ADR38800 standard; peptide; 32 AA
   Sequence 32 AA;
  Marks JD, Amersdorfer P;
   31-AUG-1998; 98US-00144886.
01-AUG-2002; 2002US-0400721P.
   09-SEP-2004.
   US2004175385-A1.
   antibacterial; antibody; botulinum neurotoxin type A; BoNT/A;
  WPI; 2004-652009/63.
   New isolated antibody that neutralizes botulinum neurotoxin type A, useful for diagnosing botulism or for treating pathologies associated
  01-AUG-2003; 2003US-00632706.
   framework 3.
   (REGC ) UNIV CALIFORNIA.
   Local Similarity
  32; Conservative
  1 RFTISRDNSKNTLFLOMNSLRADDTAVYYCAR 32
  RFTISRDNSKNTLFLQMNSLRADDTAVYYCAR
  98US-00144886.
   100.0%;
  0; Mismatches
  Score 165; DB 8;
Pred. No. 9.9e-16;
Mismatches 0;
  32
  Length 32;
  Indels
  0
  Gaps
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The invention describes an isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a specific clone where (I) binds to and neutralises botulinum neurotoxin type A

AAU02501-AAU02635, and AAU02641-AAU02748 represent the amino acid sequences of anti-adipocyte monoclonal antibody heavy chain, light and heavy chain complementarity determining regions (CDR) of the

Claim 1; Page 111; 182pp; English.

Example 4; SEQ ID NO 202; 110pp; English.

botulinum neurotoxin poisoning.

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RESULT 5
AAU02529
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   CC (BONT/A). An isolated antibody (I) that specifically binds to an epitope CC specifically bound by an antibody expressed by a clone chosen from clone CS 25, C25, C39, IC6, 3D12, B4, IP3, huC25, Ar1, Ar2, WR1(V), WR1(T), 3-1, CC comprising BONT/A neutralising epitope dare: a polypeptide (II) CC comprising BONT/A neutralising epitope having an epitope that is CC specifically bound by an antibody expressed by clones as mentioned in (I) producing (I); and a composition (III) comprising several anticc different epitope of a botulinum neurotoxin antibodies, where each antibody is specific for a CC different epitope of a botulinum neurotoxin and the combination of CC comprising (I); and a kit comprising (I) is useful for neutralising comprising (I); and a kit comprising (I) is useful for neutralising BONT/A antibody and for neutralising a botulinum neurotoxin which CC involves contacting neurotoxin with (I) in surplus, where each of (I) is specific for a different epitope of the botulinum neurotoxin and the combination of antibodies shows greater toxin neutralisation than the single antibodies in surplus. (I) is useful for neutralism CC single antibodies in surplus. (I) is useful for diagnosing the botulism or for treating pathologies associated with botulinum neurotoxin than the combination of antibodies shows greater toxin neutralisation than the combination of antibodies shows greater toxin neutralisation than the combination of antibodies shows greater toxin neutralisation than the combination of antibodies associated with botulinum neurotoxin composition or diagnosis of botulism. This is the amino acid enables rapid detection or diagnosis of botulism. This is the amino acid combined of mouse heavy chain anti-BoNT-antibody framework region 3.
   밁
  Matches
  Query Match
Best Local &
   Panel of specific binding members of antibody molecules which bind to whole adipocytes is used in the treatment of obesity and obesity related
  11-OCT-2000; 2000WO-GB003900
   WO200127279-A1.
  Homo sapiens.
  Antibody; adipocyte; heavy chain; light chain; obesity; heart disease; complementarity determining region; CDR.
  Anti-adipocyte monoclonal antibody heavy chain, FAT 19.
  29-AUG-2001
   AAU02529;
   AAU02529 standard; protein; 113
   Sequence 32
   Edwards BM,
  12-OCT-1999;
   19-APR-2001.
   (CAMB-) CAMBRIDGE ANTIBODY TECHNOLOGY.
   2001-282031/29.
DB; AAS03429.
  Similarity
   RFTISRDNSKNTLFLQMNSLRADDTAVYYCAR 32
   A,
  Conservative
   Main SH,
  (first entry
  99US-0158812P.
  100.0%; Score 165; DB 8; 100.0%; Pred. No. 9.9e-16; tive 0; Mismatches 0;
   Vaughan TJ
   B
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   Length 32;
  Indels
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  Gaps
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Matches 31
   invention. The antibodies can be used in the treatment of obesity and obesity related diseases. The antibodies can be used to deliver drugs or pro-drugs directly to the fat mass of an obese patient or the antibody can be used as a therapeutic itself. Antibodies binding specifically to adipocytes can be used to activate the immune system to destroy the cells by complement mediated lysis. The antibodies may be labeled with a detectable label such as radiolabel, fluorescent or chemical group and used in methods of diagnosis in human subjects e.g. to determine the presence of adipocyte antigen on the surface of an adipocyte to detect or determine the presence or level of adipocytes in a cell or tissue sample. The antibodies can be used as an alternative means of treatment for obese patients other than undergoing surgery to remove excess fat. Antibodies for different types of fat deposits can also be produced e.g. intra-abdominal fat associated with heart disease
  Sequence 113
  31;
   Similarity
RFTISRDNSKNTLFLQMNSLRADDTAVYYCAR 32
  Conservative
   AA;
   98.2%;
96.9%;
  Score 162; DB Pred. No. 1.1e-
1; Mismatches
   1;
   162; DB 4;
No. 1.1e-14;
  Length 113;
  0
   Gaps
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RESULT 6
ABG30486
ABG304
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  밁
  ABG30486 standard; protein; 116 AA
   anti-CD40 monoclonal antibody 12D9 heavy chain
   98
```

Human; heavy chain; CD40; autoimmune disease; multiple sclerosis; systemic lupus erythematosus; psoriasis; inflammatory bowel disease; Crohn's disease; rheumatoid arthritis; organ rejection; lymphoma; non-Hodgkin's lymphoma; monoclonal antibody; B cell; glycoprotein;

Homo sapiens.

WO200228904-A2

02-OCT-2001; 2001WO-US030857

02-OCT-2000; 2000US-0237556P

(CHIR ) CHIRON CORP.

Wang ú Yoshihara C, Donnelly JJ;

WPI; 2002-405169/43

A human anti-CD40 monoclonal antibody or fragment useful for inhibiting prollferation, growth or differentiation of a normal human B cells and treating autoimmune disease such as rheumatoid arthritis or systemic erythematosus.

Claim 8; Fig 6; 75pp; English.

The invention relates to a human monoclonal antibody or fragment capable of specifically binding to a human CD40 antigen (a glycoprotein expressed on the surface of human B cells), where the antibody or fragment is free of significant agonistic activity when it binds to the CD40 antigen, and the growth or differentiation is inhibited. The fragments comprise the complementarity determining region (CDR) of the light and heavy chains of the monoclonal antibodies secreted by a hybridoma consisting of 1588, 2024, 12D9, 9F7 and 13B4. Also included are the nucleic acids encoding

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   the antibody (or fragments). The antibodies or fragments are used for inhibiting proliferation, growth or differentiation of a normal human B cells and to inhibit antibody production by B cells. They may also be useful for treating autoimmune diseases, such as systemic lupus erythematosus, psoriasis, multiple sclerosis, inflammatory bowel disease (Crohn's disease), rheumatoid arthritis, and lymphoma (especially Non-Hodgkin's Lymphoma). The present sequence represents the antibody 12D9
   Sequence 116 AA;
 9
                1 RFTISRDNSKNTLFLQMNSLRADDTAVYYCAR 32
   Similarity
RETISRONSKNTLYLOMNSLRADDTAVYYCAR
  Conservative
   98.2%;
96.9%;
  1;
   Score 162;
Pred. No. 1
  Mismatches
   B
   2e-14;
   97
   ភ
   Length 116;
  Indels
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Human anti-CD40 monoclonal antibody 9F7 heavy
   ABG30484 standard;
                                    (first entry
   protein;
     chain
```

0

systemic lupus erythematosus; psoriasis; inflammatory bowel diseas Crohn's disease; rheumatoid arthritis; organ rejection; lymphoma; non-Hodgkin's lymphoma; monoclonal antibody; B cell; glycoprotein; Human; heavy chain; proliferation. CD40; autoimmune disease; multiple sclerosis;

Homo sapiens.

WO200228904-A2

11-APR-2002

02-OCT-2001; 2001WO-US030857.

02-OCT-2000; 2000US-0237556P

CHIRON CORP.

Wang C, Yoshihara C, Donnelly JJ;

2002-405169/43.

A human anti-CD40 monoclonal antibody or fragment useful for inhibiting proliferation, growth or differentiation of a normal human B cells and treating autoimmune disease such as rheumatoid arthritis or systemic Budul erythematosus.

Claim 6; Fig 6; 75pp; English.

RESULT 7
ABG30484
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CC Cel Inh The invention relates to a human monoclohal antibody or fragment capable of specifically binding to a human CD40 antigen (a glycoprotein expressed on the surface of human B cells), where the antibody or fragment is free of significant agonistic activity when it binds to the CD40 antigen, and the growth or differentiation is inhibited. The fragments comprise the complementarity determining region (CDR) of the light and heavy chains of the monoclonal antibodies secreted by a hybridoma consisting of 15B8, 20C4, 12D9, 9F7 and 13E4. Also included are the nucleic acids encoding the antibody (or fragments). The antibodies or fragments are used for inhibiting proliferation, growth or differentiation of a normal human B cells and to inhibit antibody production by B cells. They may also be useful for treating autoimmune diseases, such as systemic lupus erythematosus, psoriasis, multiple sclerosis, inflammatory bowel disease crythm's disease' and lumnhous (sensecially Non-

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Hodgkin's Lymphoma). heavy chain

The

present sequence represents the antibody 9F7

Sequence 117 AA;

Query Match
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Matches 31; Conserv

Conservative

1; Mismatches Score 162; DB 5; Pred. No. 1.2e-14;

0

Indels

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Gaps

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Length 117

98.2%;

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66 μ.

RETISRONSKNTLYLOMNSLRADDTAVYYCAR RETISRDNSKNTLFLOWNSLRADDTAVYYCAR 32

97

ADW08865 standard; protein; 118

B

07-APR-2005 ADW08865

(first entry)

IGF-IR antibody 22 heavy chain

variable

region,

SEQ ID

Cytostatic; Antibody; antibody therapy; antibody production; insulin-like growth factor I receptor; IGF-IR;

Homo sapiens.

Location/Qualifiers

heavy chain variable region.

Region Region

50. .66
/label=\_Complementarity\_Determining\_Region\_2\_(CDR2) 31. .35
/label= Complementarity\_Determining\_Region\_1\_(CDR1)

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ADMORBS ADMORB ADMORB ADMORB ADMORB ADMORB ADMORD AND ADMORD AND ADMORD AND ADMORD ADM
The present invention relates to antibodies 18 and 22, (A1) which bind to insulin like growth factor I receptor (IGF-IR). The antibody is capable of inhibiting the binding of IGF-I and IGF-II to IGF-IR, and is of the IGG1 isotype. The antibodies induce cell death of 20% or more cells of a preparation of IGF-IR expressing cells by antibody dependent cellular toxicity (ADCC). (A1) are useful for making a pharmaceutical composition
   Novel antibody capable of inhibiting factor I (IGF-I) and IGF-II to IGF-I
   Graus Y,
   08-JUL-2004; 2004US-00886838.
   US2005008642-A1.
   Region
   Claim 6, SEQ ID NO 3; 38pp; English
  WPI; 2005-099927/11.
   Schumacher
  10-JUL-2003;
  (VWIN/
  (GRAU/)
  (VUGT/
  (REBE/)
   (PARR/)
  (MOND/)
   KUEN/
   (KOPE/)
   KOPETZKI E.
KUENKELE K.
MUNDIGL O.
  PARREN P.
REBERS F.
   VAN DE WINKEL J.
VUGT M V.
  GRAUS
   Kopetzki
r R, Van
  2003EP-00015526
   /label= Complementarity_Determining_Region_3_(CDR3)
   ۵,
   Kuenkele
Winkel J,
  K, Mundigl 0,
Vugt MV;
  binding of insulin like growth receptor, useful for treating
   Parren P,
  Rebers
  IJ
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Query Match
Best Local Similarity
Matches 31; Conserv

Conservative

98.2%;

Score 162; DB 4; Pred. No. 1.3e-14; 1; Mismatches 0;

Length 126;

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Sequence 126 AA;

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AAG93640
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   which inhibits the binding of IGF-I and IGF-II to IGF-IR, which involves combining (A1) with a carrier. (A1) is also useful for treating a patient in need of an antitumor therapy, which involves administering (A1) alone or in combination with a cytotoxic agent, its prodrug or cytotoxic radiotherapy to the patient. The present sequence is the heavy chain variable region of antibody 22.
  11-OCT-1996;
27-JUN-1997;
10-APR-1998;
   The present invention describes an isolated Rh(D) binding protein, preferably a human antibody, (I) having an amino acid sequence comprising one of the sequences (S) given in AAG93558 to AAG93669. (I) has immunostimulant activity, and can be used as an immune system stimulant. (I) can be used in diagnostic and therapeutic medicine. The antibodies are used in diagnostics that require human antibodies instead of animal antibodies, such as determine the Rh phénotype of human red blood cells. AAH68615 to AAH68726 represent the Rh phénotype of human red blood cells. AAH68615 to AAH69369. AAG93670 to AAG93697 represent anti-Rh(D) heavy chain CDR3 amino acid sequences which are given in the exemplification of the according to the content of the content invention.
  US6255455-B1
   Human; anti-Rh(D) antibody; Rh(D) binding protein; immunostimulant;
red blood cell; Rh phenotype; diagnosis; therapeutic.
   Human anti-Rh(D) antibody clone SH54 protein sequence
  14-SEP-2001
  AAG93640;
  AAG93640 standard;
   Sequence 118 AA;
  Claim 1; Col 67; 162pp; English.
   diagnostics
therapeutic
  New isolated protein, diagnostics requiring
  N-PSDB;
   WPI; 2001-388931/41.
N-PSDB; AAH68697.
  Siegel DL
   29-JAN-1999;
   03-JUL-2001.
  Homo sapiens
   (UYPE-) UNIV
   present
   67
   Similarity
  RFTISRDNSKNTLYLQMNSLRADDTAVYYCAR
  RFTISRDNSKNTLFLOMNSLRADDTAVYYCAR 32
   Conservative
   invention
   medicine.
  (first
   PENNSYLVANIA.
   96US-0028550P.
97US-00884045.
98US-0081380P.
  99US-00240274.
  protein;
  entry)
  98.2%;
  preferably a human anti-{\tt Rh}({\tt D}) antibody a human instead of an animal antibody a
  126
  1;
   Score 162; DB 9;
Pred. No. 1.2e-14;
1; Mismatches 0
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  Length 118;
   Indels
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   ng a patient (A1) alone
   Gaps
  r use
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RESULT 10
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AC AB027
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AC AB027
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DT 12-S
DX Anti
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Huma
KW Huma
KW magi
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PP 04
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PP 11
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RBSULT 11
AAG93627
ID AAG93
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XX AAG93
XX 14-SE
DT 14-SE
XX Human
XX Human
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XW red )
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   ર્શ
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   Query Match
Best Local S
Matches 31
   11-OCT-1996;
27-JUN-1997;
10-APR-1998;
29-JAN-1999;
   The invention relates to an isolated Rh(D) binding protein. The protein can be used for magnetically activated cell sorting. The protein is useful in various diagnostic and therapeutic applications in humans, including typing of blood or blood products. The present sequence represents the amino acid sequence of a human anti-Rh(D) chain
   WPI; 2003-512273/48.
N-PSDB; ACD45361.
   Human; RH(D) binding protein; blood typing; blood product; antibody; magnetically activated cell sorting.
       Human; anti-Rh(D) antibody; Rh(D) binding protein; immunostimulant; red blood cell; Rh phenotype; diagnosis; therapeutic.
  Human anti-Rh(D) antibody clone SH10 protein sequence.
  14-SEP-2001
  AAG93627;
   AAG93627 standard; protein; 127
   Sequence 126 AA;
  Claim 4; Page 49; 187pp; English.
  New human Rh(D)-binding protein useful for various diagnostic and therapeutic applications, including typing of blood or blood products.
   Siegel DL;
  04-MAY-2001;
  27-FEB-2003
   US2003040605-A1
  Anti-Rh(D) heavy chain SH54.
  12-SEP-2003
   ABO27447;
   ABO27447 standard; protein;
   (UYPE-) UNIV
   al Similarity
31; Conserv
  67
  67
   μ.
  1
   RFTISRDNSKNTLFLQMNSLRADDTAVYYCAR 32
  RFTISRDNSKNTLYLQMNSLRADDTAVYYCAR 98
   RFTISRDNSKNTLFLQMNSLRADDTAVYYCAR 32
  RFTISRDNSKNTLYLQMNSLRADDTAVYYCAR 98
   Conservative
   PENNSYLVANIA.
  2001US-00848798
  (first
  (first entry)
   96US-0028550P.
97US-00884045.
98US-0081380P.
99US-00240274.
  entry)
  98.2%;
   126
   1,
  Score 162; DB 6;
Pred. No. 1.3e-14;
   Mismatches
   A
   A
   0
   Length 126;
  Indels
   0
   Gaps
   0
  RESULT 12
ABO27434
ID ABO27
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  ঠ
   Query Match
Best Local (
  Matches
  11-OCT-1996;
27-JUN-1997;
10-APR-1998;
         11-OCT-1996;
27-JUN-1997;
10-APR-1998;
  Human_i RH(D) binding protein; blood typing; blood product; antibody; magnetically activated cell sorting.
   Siegel DL;
  US6255455-B1
   04-MAY-2001; 2001US-00848798
  US2003040605-A1
  Homo sapiens.
   Anti-Rh(D) heavy chain SH10
   ABO27434;
  ABO27434 standard;
  29-JAN-1999;
   27-FEB-2003
   12-SEP-2003
   (UYPE-) UNIV
   83
   (first entry)
         96US-0028550P.
97US-00884045.
98US-0081380P.
```

protein; 127

A

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The present invention describes an isolated Rh(D) binding protein, preferably a human antibody, (I) having an amino acid sequence comprising one of the sequences (S) given in AAG93558 to AAG93669. (I) has immunostimulant activity, and can be used as an immune system stimulant. (I) can be used in diagnostic and therapeutic medicine. The antibodies are used in diagnostics that require human antibodies instead of animal antibodies, such as determine the Rh phenotype of human red blood cells. AAH68615 to AAH68726 represent the nucleotide sequence which encode AAG93558 to AAG93669. AAG93670 to AAG93697 represent anti-Rh(D) heavy chain CDR3 amino acid sequences which are given in the exemplification of the present invention
  New isolated protein, preferably a human anti-Rh(D) antibody for use diagnostics requiring a human instead of an animal antibody and in therapeutic medicine.
   Sequence 127 AA;
  Claim 1; Col 65; 162pp; English.
   WPI; 2001-388931/41.
N-PSDB; AAH68684.
   lomo sapiens
  / Match 98.2%;
Local Similarity 96.9%;
nes 31; Conservative
   1 RFTISRDNSKNTLFLOMNSLRADDTAVYYCAR 32
RFTISRDNSKNTLYLQMNSLRADDTAVYYCAR 99
   PENNSYLVANIA.
   97US-0028550P.
  98US-0081380P
  99US-00240274.
  1;
  Score 162; DB 4;
Pred. No. 1.3e-14;
1; Mismatches 0;
  Length 127;
   Indels
   0,
   Gaps
   ä
   0
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29-JAN-1999;

99US-00240274.

(UYPE-) UNIV

PENNSYLVANIA.

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ARBSULT 13
ADG30422
ID ADG30422
ID ADG30
XX ADG30
XX ADG30
XX GMAD
DT 26-FH
XX GMAD
XX GMAD
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XX HOMO
XX HOMO
XX HOMO
XX HOMO
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  Query Match
  can be used for magnetically activated cell sorting. The protein is useful in various diagnostic and therapeutic applications in humans, including typing of blood or blood products. The present sequence represents the amino acid sequence of a human anti-Rh(D) chain
  GMAD; VH; CDR; complementarity determining region; VL; scFv; single chain antibody; antidiabetic; type II diabetes; human; GMBC619
  ADG30422 standard;
   Sequence 127 AA;
  Claim 4, Page 48, 187pp; English.
  New human Rh\left(D\right)-binding protein useful for various diagnostic and therapeutic applications, including typing of blood or blood produced the specific produced by the s
  N-PSDB; ACD45348.
                        The invention relates to a novel antibody that specifically binds to a GMAD polypeptide comprising a first amino acid sequence that is at least 95% identical to a second amino acid sequence of a VH CDR (complementarity determining region) or VL CDR of an scrv (single chain antibody molecule). The antibody of the invention demonstrates
  Human GMBC619
   26-FEB-2004 (first entry)
   Claim 2; SEQ ID NO 55; 410pp; English.
  diagnosing,
diabetes.
   New antibody that specifically binds to GMAD diagnosing, monitoring, treating, preventing
  Baker KP,
   01-APR-2002; 2002US-0368813P
   28-MAR-2003; 2003WO-US009625
   16-OCT-2003
   WO2003085093-A2
   (HUMA-) HUMAN GENOME
  Local
   invention relates to
   sapiens.
  2003-512273/48.
   2003-804305/75
  8
   ADG30519
  Similarity
  RFTISRDNSKNTLFLQMVSLRADDTAVYYCAR 32
  RETISRONSKNTLYLOMNSLRADDTAVYYCAR
  Albert VR,
  Conservative
   activity and may be useful
  scrv protein
  protein;
  98.2%;
   SCI INC.
  Chowdhury
  an isolated Rh(D) binding
   244
   Score 162; DB 6;
Pred. No. 1.3e-14;
  Mismatches
  Ψ,
      for diagnosing, monitoring
  polypeptide, useful for or ameliorating type II
   <u>.</u>
  Length 127;
   Indels
   protein.
  The protein
  0;
  products.
   Gaps
   0
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The present invention describes a human monoclonal antibody (I) that specifically binds to tumour necrosis factor-alpha (TNPa) and comprises: (a) a heavy chain complementarity determining region 1 (CDR1) having the two fully defined 5 amino acid sequence (S1, ADP22417) or (S2, ADP22421); and (b) a light chain CDR1 having the two fully defined 11 amino acid sequence (S3, ADP22418) or (S4, ADP22424). Also described: (1) assaying (M1) the level of TNPa in a patient sample, comprising contacting with (I), and detecting the level of binding between the antibody and TNPa in the sample; (2) a composition comprising the antibody or its functional fragment and a carrier; (3) treating (M2) an animal suffering from a

Example 10; SEQ ID NO 144; 213pp; Engligh.

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   Matches
  Query Match
Best Local
  anti-TNFa antibody; anabolic; antiarteriosclerotic; antiarthritic; antibacterial; antiinflammatory; antipscriatic; antirheumatic; eating-disorder; immunomodulator; immunosuppressive; nephrotropic; neuroprotective; vasotrople; antiapoptotic; TNFa antagonist; TNF induced apoptosis; neoplastic disease; breast cancer; ovarian cancer; bladder cancer; lung cancer; glioblastoma; stomach cancer; endometrial cancer; kidney cancer; colon cancer; pancreatic cancer; prostrate cancer; immuno-mediated inflammatory disease; rheumatoid arthritis; glomerulonephritis; atherosclerosis; psoriasis; restenosis; autoimmune disease; Crohn's disease; graft-host reaction;
  septic
  ADP22238
   Sequence 244
   sequence
   treating,
   New recombinant human monoclonal antibody that specifically binds to Tumor Necrosis Factor-alpha, useful for treating neoplastic disease as cancers, or immuno-mediated inflammatory diseases such as rheumat
  02-DEC-2002; 2002US-0430729P
   02-DEC-2003; 2003WO-US038281.
  WO2004050683-A2
  Homo
   Human anti-TNFa antibody light chain variable region
  ADP22238,
   Manchulencho
   Babcook JS,
  Haak-Frendscho
   (ABGE-) ABGENIX INC.
  2004-480601/45.
   sapiens
   67
   31,
  shock; cachexia; anorexia; multiple sclerosis.
  monoclonal antibody; tumour necrosis factor-alpha; TNFa;
  Similarity
   ADP22237.
   standard; protein;
  preventing or ameliorating type II diabetes. The current is that of the human scFv protein of the invention.
  Conservative
  æ
  Kang JS, Foord O, Green L, Feng X, Kl
ho M, Rathanaswami P, Pigott C, Liang M
K, Paggioni R, Senaldi G, Qiaojuan JS;
  (first entry)
  98.2%;
  115 AA.
   ۲,
  Score
Pred.
   Mismatches
   162;
No.
  N
  DB 7;
.7e-14;
   Length
   Indels
  Klakamp
  Ĭ,
  SEQ
  诘
   0
  rheumatoid
  NO:144.
   Gaps
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RESULT 15
AD036356
AD036356
AC AD036
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  S
  CC neoplastic, or an immuno-mediated inflammatory disease by selecting an CC animal in need of treatment for the disease by administering the human CC monoclonal antibody of (1); and (4) inhibiting (M3) TWPa induced cC apoptosis in an animal by selecting an animal in need of treatment for CT TWPA induced apoptosis by administering the human monoclonal antibody of (1) (1) has anabolic, antiarteriosclerotic, antiarthritic, eating-CC (1), (1) has anabolic, antiarteriosclerotic, antiarthritic, eating-CC disorders, immunomodulator, immunosuppressive, nephrotropic, concurportective, vasctropic and antiapoptotic activities, and can be used as TWFa antagonist. The antibody (1) is useful in the preparation of CC medicament for treating TWF induced apoptosis, neoplastic disease such as tomach cancer, ovarian cancer, bladder cancer, lung cancer, glioblastoma, CC breast cancer, endometrial cancer, kidney cancer, colon cancer, CC glaesses such as rheumatoid arthritis, glomerulonephritis, CC diseases such as rheumatoid arthritis, glomerulonephritis, CC atherosclerosis, psoriasis, restenosis, autoimmune disease, Crohn's CC disease, graft-host reactions, septic shock, cachexia, anorexia, and CC multiple sclerosis. The present sequence represents a human anti-TWFa cCC antibody light chain variable region, which is used in the
   Query Match
Best Local S
Matches 31
  Determining the ability of an immunoglobulin single domain to bind to target in an intracellular environment by assessing the intracellular interaction between the immunoglobulin domain and the target by
   Unidentified
   Sequence 115 AA;
  WPI; 2004-431946/40
   Rabbitts TH, Tanaka
   15-NOV-2002; 2002GB-00026729.
  14-NOV-2003; 2003WO-GB004942.
  03-JUN-2004
  WO2004046185-A2
  immunoglobulin single domain; intracellular environment;
intracellular interaction; immunoglobulin domain; scFv;
single chain variable fragment.
  Intracellular interaction-related scFv protein SeqID20.
   26-AUG-2004 (first entry)
   ADO36356 standard; protein; 121
  (MEDI-) MEDICAL RES COUNCIL.
   66
   1 RFTISRDNSKNTLFLQMNSLRADDTAVYYCAR 32
  31;
  Similarity
   RETISRONSKNTLFLOMNSLRAEDTAVYYCAR
   Conservative
  97.6%;
96.9%;
  H
   1; Mismatches
  Score 161;
Pred. No. 1
  ₽
   DB 8;
  6e-14;
  97
   0
  Length 115;
   0
   Gaps
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This invention relates to a novel method of determining the ability of an immunoglobulin single domain to bind to a target in an intracellular environment comprising assessing the intracellular interaction between the immunoglobulin domain and the target by monitoring the signal. The

monitoring

the signal. SEQ ID

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20;

66pp; English

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                                    S
  method comprises providing a first molecule and a second molecule, where creation of the first and second molecules leads to the generation of a signal; providing a single intracellular immunoglobulin domain which is associated with the first molecule, where the single immunoglobulin domain is free of complementary immunoglobulin domains; providing an intracellular target which is associated with the second molecule, such that association of the immunoglobulin domain and the target leads to stable interaction of the first and second molecules and generation of the signal, and assessing the intracellular interaction between the immunoglobulin domain and the target by monitoring the signal. The methods are useful for determining the ability of an immunoglobulin single domain to bind to a target in an intracellular carried environment. The present sequence is that of a single chain variable fragment (scFv) protein which was used to illustrate the method of the
   Query Match
Best Local S
Matches 31
   Sequence 121 AA;
   invention.
67 RFTISRDNSKNTLFLQMNSLRAEDTAVYYCAR
                    1 RFTISRDNSKNTLFLQMNSLRADDTAVYYCAR 32
  31;
  Similarity
  Conservative
  97.6%;
  1:
  Score 161; DB 8;
Pred. No. 1.7e-14;
1; Mismatches 0
  98
  0
   Length 121;
  indels
  0
  Gaps
  0
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Search completed: November 21, Job time: 43.8425 secs 2005, 12:20:06

| 1 158 95.8 97 2 \$26935                  | Result Query No. Score Match Length DB ID Description                                                                                          | Database: PIR 80:*  1: pir1:*  2: pir2:*  3: pir3:*  4: pir4:*  Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.  SUMMARIES | Minimum DB seq length: 0 Maximum DB seq length: 2000000000 Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries | umber of hits sa | Gapop 10.0 , Gapext 0.5<br>Searched: 283416 seqs, 96216763 residues | -632-706-196<br>ISRDNSKVTLFLQMNSL                                                                                                        | Run on: November 21, 2005, 11:49:36 ; Search time 7.79487 Seconds<br>(without alignments)<br>394.995 Million cell updates/sec | GenCore version 5.1.6 Copyright (c) 1993 - 2005 Compugen Ltd. OM protein - protein search, using sw model |
|------------------------------------------|------------------------------------------------------------------------------------------------------------------------------------------------|-------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|--------------------------------------------------------------------------------------------------------------------------------------------|------------------|---------------------------------------------------------------------|------------------------------------------------------------------------------------------------------------------------------------------|-------------------------------------------------------------------------------------------------------------------------------|-----------------------------------------------------------------------------------------------------------|
| Qy 1 RPTISRDNSKNTLFLQMNSLRADDTAVYYCAR 32 | Query Match 95.8; Score 158; DB 2; Length 97; Best Local Similarity 93.8%; Pred. No. 9.6e-15; Matches 30; Conservative 2; Mismatches 0; Indels | repertoire of human germline V(H) sequen umber: \$26885; MUID:93021117; PMID:14043 \$26935 \$16935 \$100 \$100 \$100 \$100 \$100 \$100 \$100 \$10                                                                                                                             |                                                                                                                                            | RESULT 1         | ALIGNMENTS                                                          | 155 93.9 111 2 PH1643 I9<br>155 93.9 112 2 PH1647 I9<br>155 93.9 117 1 H3H026 I9<br>155 93.9 117 2 A45953 I9<br>155 93.9 117 2 A34964 I9 | 155 93.9 104 2 824255 19<br>155 93.9 104 2 824256 19<br>155 93.9 106 2 824256 19<br>155 93.9 108 2 PH1649 19                  | 93.9982                                                                                                   |

0; Gaps

0;

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Ig heavy chain V region (VH26-DN1-DXP1-JH4) - human (JSpecies: Homo sapiens (man) CJSpecies: Homo sapiens (man) CJDate: 19-Feb-194 #sequence_revision 10-Nov-1995 #text_change 30-May-1997 CJACCESSION: $24249 RJStewart, A.K.; Huang, C.; Stollar, B.D.; Schwartz, R.S. submitted to the BMBL Data Library, June 1992 AJDESCTIPTION: A single VH gene predominates in the rearranged and expressed AJREFERENCE number: $24247
  A;Cross-references: UNIPARC:UPI000002DD16; EMBL:Z17394; NID:g32843; PIDN:CAA78997.1; A;Note: designated COS-8
A;Note: designated COS-8
R;Tomilnson, I.M., Walter, G.; Marks, J.D.; Llewelyn, M.B.; Winter, G.
J. Mol. Biol. 227, 776-798, 1992
J. Mol. Biol. 227, 776-798, 1992
A;Title: The repertoire of human germline V(H) sequences reveals about fifty groups of A;Reference number: S26885; MUID:93021117; PMID:1404388
A;Accession: S26888
A;Status: preliminary
  $29546

Ig heavy chain V region (COS-8 / DP-46) - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 07-Jan-1994 #sequence_ revision 17-Nov-1995 #text_change 23-Jul-1999
C;Datesion: 829546; 826888
C;Accession: 829546; 826888
C;Tomlinson, M.; Walter, G.; Cook, G.P.; Winter, G.
submitted to the EMBL Data Library, October 1992
A;Reference number: 829543
  A;Cross-references: UNIPARC:UPI000002DD16; EMBL:Z12346; NID:g32912; PIDN:CAA78216.1; PID A;Note: designated DP-46
G;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;15-98/Domain: immunoglobulin homology <IMM>
RESULT 5
PH1642
Ig heavy chain V region
C;Species: Homo sapiens
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   #;Cross-references: UNIDARC:UDIO000176B3E; EMBL:X67070
C;Superfamily: immunoglobulin V region; immunoglobulin
C;Keywords: heteroctetramer; immunoglobulin
C;Keywords: heteroctetramer; immunoglobulin
F;10-92/Domain: immunoglobulin homology <IMM>
  밁
   S
   A;Accession: S29546
A;Molecule type: DNA
A;Residues: 1-98 <TOM>
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A;Status: prelimina
   A; Molecule type: DNA
A; Residues: 1-98 <TO2>
   A;Residues: 1-105 <STE>
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Best Local S
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  Query Match
Best Local S
Matches 30
   61
  67
  66
   1 RFTISRDNSKNTLFLQMNSLRADDTAVYYCAR 32
   1 RFTISRDNSKNTLFLOMNSLRADDTAVYYCAR 32
  30;
   30;
   95.8%;
Similarity 93.8%;
  Similarity
  RFTISRDNSKVTLYLOMNSLRAEDTAVYYCAR 98
  HILLING TO STATE THE STATE OF T
   Conservative
  Conservative
  95.8%;
93.8%;
            (clone
  5A10)
   2;
  2; Mismatches
   Score 158; DB 2;
Pred. No. 9.7e-15;
  Score 158; DB 2; Length 105; Pred. No. 1e-14;
   Mismatches
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  human (fragment)
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  97
   0
  0
   Length 98;
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   Indels
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   Gaps
  Gaps
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  human
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Ig heavy chain V region (clone 6H12) - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 24-Feb-1994 #sequence_revision 24-Feb-1994 #text_change 31-Dec-2004
C;Accession: PH1646
R;Hillson, J.L; Karr, N.S.; Oppliger, I.R.; Mannik, M.; Sasso, E.H.
J. Exp. Med. 178, 331-336, 1993
J. Exp. Med. 178, 331-336, 1993
A;Title: The structural basis of germline-encoded VH3 immunoglobulin binding to staphyloc A;Reference number: PH1642; MUID:93301610; PMID:8315388
A;Accession: PH1646
A;Molecule type: mRNA
A;Residues: 1-109 cHIL>
A;Residues: 1-109 cHIL>
A;Residues: 1-109 cHIL>
A;Residues: 1-109 cHIL>
A;Residues: 1-109 cHIL>
   C;Date: 24-Feb-1994 #sequence_revision 24-Feb-1994 #text_change 31-Dec-2004 C;Accession: PH1642 R;Hillson, J.L.; Karr, N.S.; Oppliger, I.R.; Mannik, M.; Sasso, E.H. J. Exp. Med. 178, 331-336, 1993 A;Title: The structural basis of germline-encoded VH3 immunoglobulin binding A;Reference number: PH1642; MUID:93301610; PMID:8315388 A;Accession: PH1642 A;MOID:93301610; PMID:8315388 A;Accession: PH1642 A;MOID:93301610; PMID:8315388 A;Accession: PH1642; MUID:93301610; PMID:8315388 C;Superiamily: immunoglobulin homology C;Keywords: heterotetramer; immunoglobulin F;7-90/Domain: immunoglobulin homology <IMM>
  C;Accession: PH1644
R;Hillson, J.L.; Karr, N.S.; Oppliger, I.R.; Mannik, M.; Sasso, E.H.
J. Exp. Med. 178, 331-336, 1993
A;Title: The structural basis of germline-encoded VH3 immunoglobulin binding to staphyloc A;Reference number: PH1642; MUID:93301610; PMID:8315388
A;Reference number: PH1642; MUID:93301610; PMID:8315388
A;Residues: 1-109 <- MIL-
A;Residues: 1-109 <- MIL-
A;Residues: 1-109 <- MIL-
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  Ig heavy chain V region (clone 5D11) - human (fragment)
C;Species: Homo sapiens (man)
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  A;Cross-references: UNIPROT:Q8WUK1; UNIPARC:UP10000176B7C C;Superfamily: immunoglobulin homology C;Keywords: heterotetramer; immunoglobulin F;7-90/Domain: immunoglobulin homology <IMM>
   C;Superfamily: immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
   F;7-90/Domain: immunoglobulin homology <IMM>
  PH1646
   RESULT 6
Query Match
Best Local S
Matches 30
   Query Match
Best Local S
Matches 30
   Matches
   Query Match
Best Local Similarity
                              Local Similarity
  59 RFTISRDNSKNTLYLOMNSLRAEDTAVYYCAR 90
  59
   ,
   1 RFTISRDNSKNTLFLQMNSLRADDTAVYYCAR 32
  30;
  Similarity
  RFTISRDNSKNTLFLQMNSLRADDTAVYYCAR 32
  RFTISRDNSKWTLYLQMNSLRAEDTAVYYCAR
   Conservative
  Conservative
  95.8%;
   95.8%;
95.8%; Score 158; DB 2;
93.8%; Pred. No. 1.1e-14;
tive 2; Mismatches 0
  Score 158; DB 2;
Pred. No. 1.1e-14;
2; Mismatches 0
  2
  Score 158; DB 2;
Pred. No. 1.1e-14;
   Mismatches
   DB 2; Length 108;
   90
   0
   Length 109;
   Length 109
     Indels
  Indels
     0
   0;
   0,
     Gaps
   Cape
   Gaps
   binding
   ö
     0
   0
   0
   staphyloc
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RFTISRDNSKNTLFLQMNSLRADDTAVYYCAR 32

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59

RFTISRDNSKNTLYLQMNSLRAEDTAVYYCAR

90

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A; Residues: 1-114 < FIG-
A; Cross-references: UNTC; Superfamily: 4-
C; Reyword
   R;Hillson, J.L.; Karr, N.S.; Oppliger, I.R.; Mannik, M.; Sasso, B.H.
J. Exp. Med. 178, 331-336, 1993
A;Title: The structural basis of germline-encoded VH3 immunoglobulin binding to staphyld A;Reference number: PH1642; MUID:93301610; PMID:8315388
A;Accession: PH1645
   R;Figini, M.; Marks, J.D.; Winter, G.; Griffiths, A.D. J. Mol. Biol. 239, 68-78, 1994
A;Title: In vitro assembly of repertoires of antibody chains A;Reference number: S46390; MUID:94254092; PMID:8196048
A;Accession: S46390
   A; Molecule type: mRNA
A; Residues: 1-111 <HIL>
A; Residues: UNIPROT: QBWUK1; UNIPARC: UPI0000176B7B
A; Cross-references: UNIPROT: QBWUK1; UNIPARC: UPI0000176B7B
C; Superfamily: immunoglobulin
C; Keywords: heterotetramer; immunoglobulin
   Ig heavy chain V region - human
C;Species: Homo sapiens (man)
C;Date: 27-Jan-1995 #sequence_revision 27-Jan-1995 #text_change 20-Jun-2000
C;Accession: S46390
  Ig heavy chain V region - human

(;Species: Homo sapiens (man)

(;Date: 27-Jan-1995 #sequence_revision 27-Jan-1995 #text_chan

(;Accession: $46391

R;Pigini, M.; Marks, J.D.; Winter, G.; Griffiths, A.D.

J. Mol. Biol. 239, 68-78, 1994

A;Title: In vitro assembly of repertoires of antibody chains

A;Reference number: $46390; MUID:94254092; PMID:8196048

A;Accession: $46391
  밁
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   á
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-114 <FIG>
  RESULT 10
  S
   g heavy chain V region (clone 6C8) - human (fragment);Species: Homo sapiens (man);Date: 24-Feb-1994 #sequence_revision 24-Feb-1994 #te;Accession. DM'45E
   7-90/Domain: immunoglobulin homology <IMM>
  Accession: PH1645
  ;ross-references: UNIPARC:UPI000011663F; EMBL:Z31686; NID:g509782; Superfamily: immunoglobulin V region; immunoglobulin homology ;Keywords: heterotetramer; immunoglobulin ;15-98/Domain: immunoglobulin homology <IMM>
  Matches
   Matches
   Local Similarity
  ocal Similarity
  59
   67
  30;
   30;
   RFTISRDNSKNTLFLQMNSLRADDTAVYYCAR 32
  RFTISRDNSKNTLFLQMNSLRADDTAVYYCAR 32
  RETISRONSKNTLYLOMNSLRAEDTAVYYCAR
   Conservative
   Conservative
  95.8%;
   95.8%;
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   2;
  Score 158; DB 2;
Pred. No. 1.1e-14;
2; Mismatches 0
   Score 158; DB 2;
Pred. No. 1.1e-14;
2; Mismatches 0
  24-Feb-1994 #text_change 31-Dec-2004
   27-Jan-1995 #text_change
  Length 114;
   Length 111;
  Indels
   Indels
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   on the
  the
   20-Jun-2000
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  surface of phage
   surface of phage
  PIDN: CAA83491.1;
   Gaps
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85

RPTISRDNSKNTLYLQMNSLRAEDTAVYYCAR 116

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RESULT 11
S46392
   A;Cross-references: UNIPARC:UPI00001137D5; EMBL:Z31687; NID:g509784; PIDN:CAA83492.1; PII C;Superfamily: immunoglobulin V region; immunoglobulin homology C;Keywords: heterotetramer; immunoglobulin F;15-98/Domain: immunoglobulin homology <IMM>
  Ig heavy chain precursor V-region chain 5A - human C;Species: Homo sapiens (man) C;Species: Homo sapiens (man) C;Species: 30-Jun-1999 #sequence_revision 30-Jun-1989 #text_change 16-Aug-1996 C;Accession: B28966; A32782 R;Wilson, M.R.; Middleton, D.; Warr, G.W. R;Wilson, M.R.; Middleton, D.; Warr, G.W. A;Title: Immunoglobulin heavy chain variable region gene evolution: structur A;Reference number: A28966; MUID:88144476; PMID:3125551 A;Accession: B28966
  R;Figini, M.; Marks, J.D.; Winter, G.; Griffiths, A.D. J. Mol. Biol. 239, 68-78, 1994
A;Title: In vitro assembly of repertoires of antibody chains A;Reference number: S46390; MUID:94254092; PMID:8196048
A;Accession: S46392
   A;Residues: 1-114 <FIG>A;Residues: 1-114 <FIG>A;Residues: 1-114 <FIG>B;Residues: 1-114 <FIG>B;Residues: NID:g499306; PIDN:CAA83493.1; PII A;Cross-references: UNIPARC:UPI00001137D6; EMBL:Z31688; NID:g499306; PIDN:CAA83493.1; PII C;Superfamily: immunoglobulin V region; immunoglobulin homology C;Keywords: heterotetramer; immunoglobulin C;Keywords: heterotetramer; immunoglobulin homology <IMM**
   Ig heavy chain V region (VH-28) - human
C;Species: Homo sapiens (man)
C;Date: 27-Jan-1995 #sequence_revision 27-Jan-1995 #text_change
   밁
   5
  A;Note: the authors note that this sequence is of higher primate (probably human) origin C;Superfamily: immunoglobulin V region; immunoglobulin homology C;Keywords: heterotetramer; immunoglobulin
   A;Cross-references: UNIPARC:UPI000012CEDE R;Wilson, M.R.; Middleton, D.; Warr, G.W. Proc. Natl. Acad. Sci. U.S.A. 86, 3276, 1989
  밁
  á
S
  A;Reference number: A32782
A;Contents: annotation; erratum
  B28966
  A;Status: preliminary A;Molecule type: DNA
   A;Molecule type: DNA
A;Residues: 1-116 <WIL>
  RESULT 12
  Accession: S46392
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  Query Match
Best Local Similarity
   Matches
  34-116/Domain: immunoglobulin homology <IMM>
  Query Match
  Matches
  Query Match
Best Local
   Local
  67
   67
  30;
   30;
  Similarity
  Similarity
  RFTISRDNSKNTLFLQMNSLRADDTAVYYCAR 32
  RFTISRDNSKNTLYLQMNSLRAEDTAVYYCAR
  RFTISRDNSKNTLFLOMNSLRADDTAVYYCAR
    RFTISRDNSKNTLFLOMNSLRADDTAVYYCAR 32
  Conservative
   Conservative
  95.8%;
  95.8%;
   95.8%;
  Score 158; DB 2;
Pred. No. 1.1e-14;
2; Mismatches 0
   Score 158; DB 2;
Pred. No. 1.1e-14;
2; Mismatches 0
  <u>ب</u>
  Score 158; DB 2;
Pred. No. 1.2e-14;
2; Mismatches 0
  98
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   Length
  Length 114;
   Length 116;
   Indels
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   of phage by
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   and
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Ig heavy chain V region (M74) - human
(7)Species: Homo sapiens (man)
(7)Date: 21-Dec-1990 #sequence_revision 21-Dec-1990 #text_change 31-Dec-2004
(7)Accession: G36005
R;Schroeder Jr., H.W.; Wang, J.Y.
Proc. Natl. Acad. Sci. U.S.A. 87, 6146-6150, 1990
A;Title: Preferential Sci. U.S.A. 87, 6146-6150, 1990
A;Reference number: A36005; MUID:90349571; PMID:2117273
A;Accession: G36005
   A;Molecule type: mRNA
A;Residues: 1-119 <RAA>
A;Residues: 1-119 <RAA>
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C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;15-98/Domain: immunoglobulin homology <IMM>
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C;Species: Homo sapiens (man)
C;Date: 02-Dec-1993 #sequence_revision 26-May-1995 #text_change 17-Mar-1999
C;Accession: 831111
  A;Map position: 14q32.33-14q32.33
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C;Keywords: heterotetramer; immunoglobulin
F;15-98/Domain: immunoglobulin homology <IMM>
   C;Accession: F36005

C;Accession: F36005

C;Accession: F36005

C;Accession: F36005

C;Accession: F36005

C;Accession: F36005

C;Accession: F36005

C;Accession: F36005
  Ig heavy chain V region (M49) - human
C;Species: Homo sapiens (man)
C;Date: 21-Dec-1990 #sequence_revision 21-Dec-1990 #text_change 31-Dec-2004
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  श्
  C;Genetics:
A;Gene: GDB:IGH@; IGHDY1
  A; Molecule type: mRNA
A; Residues: 1-119 < SCH>
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   5
  A;Cross-references: UNIPROT:QBWUK1; UNIPARC:UPI0000176C32; GB:M34026
   Status: preliminary
  Query Match
   Cross-references: GDB:118731; OMIM:146910
  Matches
  Query Match
Best Local
   Matches
   y Match 95.8%;
Local Similarity 93.8%;
hes 30; Conservative ;
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  67
  1 RFT1SRDNSKNTLFLQMNSLRADDTAVYYCAR 32
  1 RFTISRDNSKVTLFLQMNSLRADDTAVYYCAR 32
  30,
  Similarity 93.8
30; Conservative
  RFTISRDNSKNTLYLOMNSLRAEDTAVYYCAR 98
  95.8%;
   Score 158; DB 2;
Pred. No. 1.2e-14;
2; Mismatches 0
  2
  Score 158; DB 2;
Pred. No. 1.2e-14;
2; Mismatches (
  Length 119;
  Length 119,
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  Indels
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  0,
  Gaps
  0
   0
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A;Gene: GDB:IGH@; IGHDY1
A;Cross-references: GDB:118731; OMIM:146910
A;Map position: 14q32.33-14q32.33
C;Superfamily: immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;15-98/Domain: immunoglobulin homology <IMM>
  A;Status: preliminary
A;Molecule type: mRNA
A;Mesidues: 1-121 < <CH>
A;Cross-references: UNIPROT:Q8WUK1; UNIPARC:UPI0000176C2C; GB:M34031
C;Genetics: CTIONIC C;Geneti
   δ
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   1 RETISRONSKNTLFLOMNSLRADDTAVYYCAR 32
  5:81; 93.81; 93.81;
RFTISRDNSKNTLYLQMNSLRAEDTAVYYCAR
   Conservative
   2; Mismatches
  Score 158; DB 2;
Pred. No. 1.2e-14;
        98
   Length 121;
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Search completed: November 21, 2005, 12:22:17 Job time: 7.79487 secs

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Result
No.
  Title:
Perfect score:
   Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
   Minimum DB seq length: 0
Maximum DB seq length: 200000000
   Sequence:
   OM protein -
  Total number of hits satisfying chosen parameters:
   Scoring table:
   Database
   8
   Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
  Score
   protein search, using sw model
Query
Match
  BLOSUM62 .
Gapop 10.0 , Gapext 0.5
   November 21, 2005, 11:45:55; Search time 47.3553 Seconds (without alignments)
476.756 Million cell updates/sec
  2166443 seqs, 705528306 residues
   1 RFTISRDNSKNTLFLQMNSLRADDTAVYYCAR 32
   UniProt_05.80:*
1: uniprot_sprot:*
2: uniprot_trembl:*
   US-10-632-706-196
  GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
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| RESULT HV05_CC PD DT 00 DT 00 DT 11 DT 11 DT 11 COC CC CC CC RN [                                                                                                                                                                                                                                          | ል 5<br>8 8                                                                                 | SO SO SO SO SO SO SO SO SO SO SO SO SO S                                                                                                                            | RESULT Q9UGP3 ID Q DT 0 DT 0 DT 0 DT 0 OC DT 0 OC C C C C C C C C C C C C C C C C C C                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          |                                                                                                                                                                                             |
|------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|--------------------------------------------------------------------------------------------|---------------------------------------------------------------------------------------------------------------------------------------------------------------------|--------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|---------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| ULT 2  CARAU HV05 P1918 01-NO 01-NO 10-MA IG he Caras Eukar Actin Cypri NCBI [1]                                                                                                                                                                                                                           | Best Local Best Local Matches  7                                                           | AUCLEOTIDE Zafiropoul Kramboviti Kramboviti Submitted EMBL; AJ13 HSSP; P017 SMR; O9UGP Ensembl; ElinterPro; InterPro; SMART; SMO PROSITE; PNON TER NON TER SEQUENCE | T 1 3 HUM 9 UGP Q9 UGP Q9 UGP Q0 1 - MA 01 - MA 01 - MA 01 - MA 01 - MA Mamma Homo Eukar Mamma Mamma                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           | 00000000000000000000000000000000000000                                                                                                                                                      |
| CARAU STJ P19181; 01-NOV-1990 (Rel. 01-NOV-1990 (Rel. 10-NAY-2005 (Rel. 10-MAY-2005 (Rel. 12 heavy chain V J Carassius auratus Eukaryota; Metazoo Actinopterygii; NCBI_TaxID=7957; [1]                                                                                                                     | 0 1 36                                                                                     | NUCLEOTIDE SEQUENTION AND AND AND AND AND AND AND AND AND AN                                                                                                        |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                | 144<br>144<br>144<br>144<br>144<br>144<br>144<br>143<br>143<br>143                                                                                                                          |
| ULT 2 5_CARAU STANDARD; P P19181; 01_NOV-1990 (Rel. 16, Created) 01_NOV-1990 (Rel. 47, Last sequ 10_MAY-2005 (Rel. 47, Last anno 1g heavy chain v region 5A prec Carassius auratus (Goldfish). Eukaryota; Metazoa; Chordata; C Actinopterygii, Neopterygii; Te Cyprinidae; Carassius. NCBI_TaxID=7957; [1] | Conserva<br>Conserva<br>FISRDNSKN                                                          | 3QUEN<br>3.;<br>3.;<br>1.19<br>50; 11<br>7.11<br>1.10<br>1.10<br>1.11<br>1.12<br>1.12<br>1.12<br>1.12                                                               | AN PRELIMINARY  0 (TrembLrel. 0 (TrembLrel. 10 (TrembLrel. 11 (TrembLrel. 12 (TrembLrel. 13 (Tremblarel. 14 (TrembLrel. 15 (Tremblarel. 16 (Tremblarel. 16 (Tremblarel. 17 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tremblarel. 18 (Tr | 87.3<br>87.3<br>887.3<br>887.3<br>887.3<br>886.7<br>886.7<br>886.1                                                                                                                          |
| TANDARD; . 16, Create . 16, Last s . 47, Last s . 47, Last s . Goldfish .s (Goldfish) .oa, Chordate Neopterygii                                                                                                                                                                                            | 93. Pative ONTLFL                                                                          | 11c<br>1443<br>11g-<br>11g-<br>11c-<br>11c-<br>11c-<br>11c-<br>11c-<br>11c-                                                                                         | ~ < K K K H                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    | 573<br>1121<br>1136<br>1136<br>1136<br>493<br>1120<br>1126<br>487<br>1126<br>1126<br>1126<br>1126                                                                                           |
| GLEST E                                                                                                                                                                                                                                                                                                    | OMENS                                                                                      | ogian<br>the ;<br>1.1;<br>1.1;<br>Ho<br>like<br>v.<br>CE; 1                                                                                                         | ;<br>13,<br>13,<br>25,<br>in (<br>data                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         | H H M M M H H M M M H H M M M M M M M M                                                                                                                                                     |
| RT;<br>ence<br>tati<br>urso<br>urso<br>leos                                                                                                                                                                                                                                                                | imilarity 93.8%; Pred. No. 4. ; Conservative 2; Mismatche RFTISRDNSKNTLFLOMNSLRADDTAVYYCAR | M.,<br>/GenB<br>RNA.<br>apien                                                                                                                                       | NARY; PRT; 11 el. 13, Created) el. 13, Last sequenc el. 25, Last annotat chain (Fragment)                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      | OBWUJB HUMAN HV3T HUMAN HV3T HUMAN HV36 HUMAN O56617 MOUSB O6007 MOUSB O8054 HUMAN HV3E HUMAN HV3E HUMAN O6007 HUMAN O6007 HUMAN O90LB6 HUMAN O90LB6 HUMAN HV3E HUMAN HV3E HUMAN ALIGNMENTS |
| llé AA.<br>odáte)<br>update)<br>update)<br>i, Vertebrata;                                                                                                                                                                                                                                                  | 7e-15;<br>8 0;<br>32                                                                       | databa<br>databa<br>E3 CRC6                                                                                                                                         | 112 AA.<br>nce update)<br>ation update)<br>; Vertebrata;<br>primates; Cata                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     |                                                                                                                                                                                             |
| Euteleostomi;; Cypriniforme                                                                                                                                                                                                                                                                                | 18                                                                                         | Borrebaeck<br>BeB.<br>BeB.                                                                                                                                          | <b></b>                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        | Q8wu38<br>P01781<br>P017781<br>P017783<br>Q6n089<br>Q6n65k4<br>P01766<br>P01772<br>Q6n093<br>Q9u1b6<br>Q6n094<br>P01763<br>P01763                                                           |
| itomi,                                                                                                                                                                                                                                                                                                     | 0;                                                                                         | O                                                                                                                                                                   | tomi;<br>Hominidae;                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            | home home home home home home home home                                                                                                                                                     |
| ies;                                                                                                                                                                                                                                                                                                       | Gарв О                                                                                     | .A.K.,                                                                                                                                                              | idae;                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          | sapien sapien sapien musculu sapien musculu sapien sapien sapien sapien sapien sapien sapien sapien sapien sapien sapien sapien sapien sapien sapien                                        |
|                                                                                                                                                                                                                                                                                                            | Ş                                                                                          |                                                                                                                                                                     |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                |                                                                                                                                                                                             |

87.995.55.55.57.79.995.55.68

Q9UGP3 HUMAN
HV05 CARAU
Q65279 HUMAN
Q61N78 HUMAN
Q61N78 HUMAN
Q9UL90 HUMAN
Q9UL72 HUMAN
Q9UL72 HUMAN
Q9UL71 HUMAN
Q9UL91 HUMAN
Q9UL91 HUMAN
Q9UL93 HUMAN
Q6RY2 HUMAN
Q6RY2 HUMAN
Q6RY2 HUMAN
Q6RY2 HUMAN
Q6RY2 HUMAN
Q6RO36 HUMAN
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Q9ugp3 P19181 Q65zc9 Q61n78

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Q6P6C4\_HUMAN

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NUCLEOTIDE SEQUENCE.
MEDLINE=88144476; PubMed=3125551;
Wilson M.R., Middleton D., Warr G.W.;
"Immunoglobulin heavy chain variable region gene evolution:

and family relationships of two genes and fish.";

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RESULT 1

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ID Q65Z5

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InterPro; IPR007110; Ig-like.
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between the Swiss Institute of Bioinformatics and the EMBL
the European Bioinformatics Institute. There are no restrict
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   "Complement recruitment using bispecific Nat. Biotechnol. 15:629-631(1997).
   Kontermann R.B., Wing M.G., Winter G.
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116
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IG_LIKB; 1.
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49
54
68
116
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Pred. No. 4.
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Framework-3.
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Framework-1.

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  Gaps
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93.8%;
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   53EB0BCEDE81076E CRC64;
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  complementarity-determining regions (CDR3) in human fetal B lymphocyte immunoglobulin heavy chain rearrangements.";
ENUT. J. Immunol. 22:247-251 (1992).
   fetus.";
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   NUCLEOTIDE SEQUENCE.
MEDLINE=98277139; PubMed=9614934; DOI=10.1006/clin.1998.4531;
MEDLINE=98277139; PubMed=9614934; DOI=10.1006/clin.1998.4531;
MAI X . Liu B., Van der Merwe P.L., Kalis N.N., Berney S.M.,
  HV3C_HUN
P01764;
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EMBL; APO35024; AND56260.1;

PIR; S78486; S78486.

HSSP; P01772; 2FB4.

SMR; Q9UL90; 1-113.
  Raaphorst F.M., Timmers B., Kenter M.J.,
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13-SEP-2005 (Rel. 48, Last annotation update)
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Eukaryota, Metazos, Chordata, Craniata, Verte
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InterPro; IPR003596; Ig_v.
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MEDLINE=81101090; PubMed=6450418;
MALTHYSSENS G., Rabbitts T.H.;
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Toyonaga B., Yoshikai Y., Vadasz V., Chin B., Mak T.W.;
Toyonaga B., Yoshikai Y., Vadasz V., Chin B., Mak T.W.;
"Organization and sequences of the diversity, joining, and const
region genes of the human T-cell receptor beta chain.";
Proc. Natl. Acad. Sci. U.S.A. 82:8624-8628(1985).
-I- SIMILARITY: Contains 1 Ig-like (immunoglobulin-like) domain.
   This Swiss-Prot entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its
  Mariette X., Tsapis A., Brouet J.C.;
"Nucleotidic sequence analysis of the variable domains of four
monoclonal IgM with an antibody activity to myelin-associated
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  glycoprotein."
NUCLEOTIDE SEQUENCE.
TISSUE=Primary B-Cells;
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MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
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RA Rodriguez A.C., Schein J.E., Jones S.J.M., Marra M.A.,

"Generation and initial analysis of more than 15,000 full-length human and mouse cDNA sequences.";
  PubMed=2117273;

Schroeder H.W. Jr, Wang J.Y.;

Schroeder H.W. Jr, Wang J.Y.;

"Preferential utilization of conserved immunoglobulin "Preferential utilization of the liment of the life.";

variable gene segments during human fetal life.";

Proc. Natl. Acad. Sci. U.S.A. 87:6146-6150(1990).
  NUCLEOTIDE SEQUENCE.

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PubMed=2538551; DOI=10.1084/jem.169.4.1391;

Nickerson K.G., Berman J., Glickman B., Chess L., Alt F.W.;

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Nickerson K.G., Berman J., Exp. Hots T., Alt F.W.;

Nickerson K.G., Alt F
  PubMed=1383695; DOI=10.1016/0161-5890(92)90173-U; Cuisinier A.M., Pumoux F., Fougereau M., Tonnelle C.; "IGM kappa/lambda EBV human B cell clone: an early stidifferentiation of fetal B cells or a distinct B line. Mol. Immunol. 29:1363-1373(1992).
   PubMed=2840480; DOI=10.1084/jem.168.1.229; Bird J., Gallli N., Link M., Stites D., Sklar J.; Continuing rearrangement but absence of somatic hypermutation immunoglobulin genes of human B cell precursor leukemia."; J. Exp. Med. 168:229-245(1988).
  Neale G.A., Kitchingman G.R.;

"mRNA transcripts initiating within the human immunoglobulin mu
"manacer region contain a non-translatable exon and are
extremely heterogeneous at the 5' end.";

Nucleic Acids Res. 19:2427-2433(1991).
   NUCLEOTIDE SEQUENCE.
MEDLINE=93301610; PubMed=8315388;
Hillson J.L., Karr N.S., Oppliger
   "Restricted utilization of germ-line VH3 genes and short diverse third complementarity-determining regions (CDR3) in human fetal B lymphocyte immunoglobulin heavy chain rearrangements.", Eur. J. Immunol. 22:247-251(1992).
   PubMed=1904154;
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  Raaphorst F.M., Timmers
  PubMed=1730252;
   NIH MGC Project;
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I.R., Mannik M., Sasso E.H
  clone: an early step of a distinct B lineage?";
  Van
   Tol M.J.,
  databases
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            E.H.;
  J.M.,
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PIR; |
PIR;
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Q9UL72;
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Makiya R., Stigbrand T.;
"Placental alkaline phosphatase has a binding silimunoglobulin-G Fc portion.";
Eur. J. Biochem. 205;341-345(1992).
EMBL; AF035042; AAD56278.1; -; mRNA.
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InterPro; IPR007110; Ig-like.
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01.MAY-2000 (TrEMBLrel. 13, Last sequence update)
01.CCT-2003 (TrEMBLrel. 25, Last annotation updat)
Myosin-reactive immunoglobulin heavy chain varial
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  EXP. Med. 178:331-336(1993).

EXP. Med. 178:331-336(1993).

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R; F36005; F36005.

R; G36005, F36005.

R; PH1642; PH1642.

R; PH1643; PH1643.

R; PH1644; PH1645.

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N.N., Berney S.M.,
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   on update)
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Submitted (OCT-2000) to the
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01-MAR-2001 (TrEMBLrel. 16,
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RA Hopkins R.F., Jordan H., Moore T., Max 9.I., Wang J., Hsieh F.,

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  SO DE RESTE 
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Last annotation update)

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  "Molecular characteristics of antibodies associated idiotype.";
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RX Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

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87.5%;
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Last annotation update)
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  Makiya R., Stigbrand T.;
"Placental alkaline phosphatase has a binding site for the human immunoglobulin-G Fc portion.";
Eur. J. Biochem. 205:341-345(1992).
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  "Complete sequencing cDNAs.";
  Okumura K., Nagase
Nakai K., Yada T.,
  PIR; S21205; S21205.
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  MEDLINE=93301610; PubMed=8315388; DOI=10.1084/jem.178.1.331; Hillson J.L., Karr N.S., Oppliger I.R., Mannik M., Sasso E.H.; "The structural basis of germline-encoded VH3 immunoglobulin binding to staphylococcal protein A.";
   MEDLINE=98277139; PubMed=9614934; DOI=10.1006/clin.1998
Wu X., Liu B., Van der Merwe P.L., Kalid N.N., Berney S.
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US-09-848-798-152
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| US-09-848-798-152 US-09-848-798-152 US-09-848-798-152 Sequence 152, Application US/09848798 Patent No. 6858719 GENERAL INFORMATION: APPLICANT: Siegel, Donald L. TITLE OF INVENTION: Rh(D)-BINDING PROTEINS AND MAGNETICALLY ACTIVATED CELL TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF FILE REPERENCE: 09596-42U2 CURRENT APPLICATION NUMBER: US/09/848,798 CURRENT FILING DATE: 2001-05-04 PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 09/240,274 PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 60/028,550 PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 60/028,550 PRIOR APPLICATION DATE: EARLIER APPLICATION NUMBER: 60/028,550 PRIOR APPLICATION DATE: EARLIER FILING DATE: 1996-10-11 | Query Match 98.2%; Score 162; DB 2; Length 126;  Best Local Similarity 96.9%; Pred. No. 1e-15;  Matches 31; Conservative 1; Mismatches 0; Indels 0; Gaps 0;  Qy 1 RFTISRDNSKNTLFLQMNSLRADDTAVYYCAR 32 | EARLIER APPLICATION NUMBER: 60/028,550 ; EARLIER PILING DATE: 1996-10-11 ; NUMBER OF SEQ ID NOS: 224 ; SOFTWARE: Patentin Ver. 2.0 ; SEQ ID NO 152 ; SEQ ID NO 156 ; TYPE: PRT ; ORGANISM: Homo sapiens ; FEATURE: ; OTHER INFORMATION: anti-Rh(D) antibody clone SH54 US-09-240-274-152 | 174-152 152, Application US/0924027 152, Application US/0924027 1625455 10000000000000000000000000000000000 | ALIGNMENTS | 28 158 95.8 98 2 US-10-194-9†5-30 Sequence 30, Appl 158 95.8 98 2 US-09-534-717-624 Sequence 624, Appl 30 158 95.8 98 2 US-09-534-717-625 Sequence 625, Appl 31 158 95.8 98 2 US-09-534-717-627 Sequence 627, Appl 32 158 95.8 98 2 US-09-534-717-629 Sequence 627, Appl 33 158 95.8 98 2 US-09-534-717-630 Sequence 628, Appl 35 158 95.8 98 2 US-09-534-717-630 Sequence 630, Appl 35 158 95.8 98 2 US-09-534-717-631 Sequence 631, Appl 36 158 95.8 98 2 US-09-534-717-632 Sequence 631, Appl 37 158 95.8 98 2 US-09-534-717-634 Sequence 634, Appl 38 158 95.8 98 2 US-09-534-717-634 Sequence 634, Appl 38 158 95.8 98 2 US-09-534-717-634 Sequence 634, Appl 39 158 95.8 98 2 US-09-534-717-641 Sequence 642, Appl 31 158 95.8 98 2 US-09-534-717-642 Sequence 643, Appl 31 158 95.8 98 2 US-09-534-717-643 Sequence 643, Appl 35 158 95.8 98 2 US-09-534-717-643 Sequence 644, Appl 35 158 95.8 98 2 US-09-534-717-643 Sequence 644, Appl 35 158 95.8 98 2 US-09-534-717-643 Sequence 644, Appl 35 158 95.8 98 2 US-09-534-717-644 Sequence 644, Appl 35 158 95.8 98 2 US-09-534-717-645 Sequence 647, Appl 35 158 95.8 98 2 US-09-534-717-645 Sequence 647, Appl 35 158 95.8 98 2 US-09-534-717-645 Sequence 647, Appl 35 158 95.8 98 2 US-09-534-717-645 Sequence 647, Appl 35 158 95.8 98 2 US-09-534-717-647 Sequence 647, Appl 35 158 95.8 98 2 US-09-534-717-647 Sequence 647, Appl 35 158 95.8 98 2 US-09-534-717-647 Sequence 647, Appl 35 158 95.8 98 2 US-09-534-717-647 Sequence 647, Appl 35 158 95.8 98 2 US-09-534-717-647 Sequence 647, Appl 35 158 95.8 98 2 US-09-534-717-647 Sequence 647, Appl 35 158 95.8 98 2 US-09-534-717-647 Sequence 647, Appl 35 158 95.8 98 2 US-09-534-717-647 Sequence 647, Appl 35 158 95.8 98 2 US-09-534-717-647 Sequence 647, Appl 35 158 95.8 98 2 US-09-534-717-647 Sequence 647, Appl 35 158 95.8 98 2 US-09-534-717-647 Sequence 647, Appl 35 158 95.8 98 2 US-09-534-717-647 Sequence 647, Appl 35 158 95.8 98 2 US-09-534-717-648 Sequence 647, Appl 35 158 95.8 98 2 US-09-534-717-648 Sequence 647, Appl 35 158 95.8 98 2 US-09-534-717-648 Sequence 647, Appl |
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GENERAL INFORMATION:

APPLICANT: Siegel, Donald L.

TITLE OF INVENTION: Rh(D)-BINDING PROTEINS AND MAGNETICALLY ACTIVATED CELL

TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF

FILE REFERENCE: 09596-42012

CURRENT APPLICATION NUMBER: US/09/240,274

CURRENT FILING DATE: 1999-01-9

EARLIER APPLICATION NUMBER: 60/081,380

EARLIER PILING DATE: 1998-04-10

EARLIER PILING DATE: 1998-04-10

EARLIER APPLICATION NUMBER: 60/028,550

EARLIER FILING DATE: 1996-10-11

NUMBER OF SEQ ID NOS: 224

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Patent No. 6888719

GENERAL INFORMATION:
APPLICANT: Siegel, Donald L.
TITLE OF INVENTION: Rh(D)-BINDING PROTEINS AND MAGNETICALLY ACTIVATED CELL
TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF
FILE REFERENCE: 09596-42U2
CURRENT APPLICATION NUMBER: US/09/848,798
CURRENT FILING DATE: 2001-05-04
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 09/240,274
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APPLICANT: GILMAN, STEVEN C.

APPLICANT: DAVIS, C. GEOFFREY

APPLICANT: CORVALAN, JOSE R.

TITLE OF INVENTION: HUMAN MONOCLONAL ANTIBODIES TO FILE REFERENCE: ARX-PF1

CURRENT APPLICATION NUMBER: US/09/472,087

CURRENT FILING DATE: 1999-12-23

PRIOR APPLICATION NUMBER: 60/113,647

PRIOR FILING DATE: 1998-12-23
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APPLICANT: GILMAN, STEVEN C.
APPLICANT: DAVIS, C. GEOFFREY
APPLICANT: CORVALAN, JOSE R.
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APPLICANT: HANKE, JEFFREY H.
APPLICANT: GILMAN, STEVEN C.
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APPLICANT: NEVEU, MARK J.
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APPLICANT: GILMAN, STEVEN C.
APPLICANT: DAVIS, C. GEOFFREY
APPLICANT: CORVALAN, JOSE R.
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APPLICANT: MUELLER, BILLEN B
APPLICANT: HANKE, JEFFREY H.
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US-08-471-780C-82
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APPLICANT: Casterman, Cecile
APPLICANT: Hamers, Raymond
TITLE OF INVENTION: Immunoglv
   TELEFAX: 202-408-4400
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
GENERAL INFORMATION:
APPLICANT: Casterman, Cecile
APPLICANT: Hamers, Raymond
TITLE OF INVENTION: Immunogle
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TYPE: amino acid
STRANDEDNESS: single
TOPOLCOY: linear
MOLECULE TYPE: protein
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COMPUTER: IBM PC compatible
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  FILING DATE: 21-AUG-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: FR 9:
FILING DATE: 21-MAY-1993
ATTORNEY/AGENT INFORMATION:
   SOPTWARE: Patentin Release #1.0, Vocurrent Application DATA:
APPLICATION NUMBER: US/08/471,780C FILING DATE: 06-UN-1995
   CLASSIFICATION: 530
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  B: Finnegan, Henderson, Farabow, Garrett & Dunner
1300 I Street, N.W.
  202-408-4400
  17-AUG-1993
17-BER: FR 92402326.0
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 Immunoglobulins Devoid of Light Chains
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   FR 93401310.3
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CORRESPONDENCE ADDRESS: NUMBER OF SEQUENCES:

130

ADDRESSEE:

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FILING DATE: 17-AUG-1993
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SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
   APPLICANT: Casterman, Cecile
APPLICANT: Hamers, Raymond
TITLE OF INVENTION: Immunoglobulins Devoid of Light Chains
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ADDRESSEE: Finnegan, Henderson, Farabow, Garrett & Dunner
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LENGTH: 32 amino acids
TYPE: amino acid
  FILING DATE: 21-MAY-1993
ATTORNEY/AGENT INFORMATION:
NAME: POLICE, Jane E.R.
REGISTRATION NUMBER: 33,332
REFERENCE/DOCKET NUMBER: 04958.0008-00000
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   06-JUN-1995
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US/08/471,282A
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APPLICATION NUMBER: FR 924
FILING DATE: 21-AUG-1992
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FILING DATE: 21-AUG-1992
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FILING DATE: 21-MAY-1993
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APPLICATION NUMBER: US/08/466,710C
                 ATTORNEY/AGENT INFORMATION:
NAME: POLICE, Jane E. R.
REGISTRATION NUMBER: 33,332
REFERENCE/DOCKET NUMBER: 04:
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REFERENCE/DOCKET NUMBER: 04
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APPLICANT: Hamers, Raymond
TITLE OF INVENTION: Immunogl
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   D.C
  Potter, Jane E.R
  s: Finnegan, Henderson, Farabow, Garrett & Dunner
1300 I Street, N.W.
  linear
  202-408-4000
   JMBER: FR 93401310.3
21-MAY-1993
   single
  Immunoglobulins Devbid of Light Chains
   FR 92402326.0
   US/08/106,944
   FR 92402326.0
  FR 93401310.3
  US 08/106,944
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   04958.0008-00000
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                   04958.0008-00000
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TELEFAX: 202-408-4400 INFORMATION FOR SEQ ID NO:

202-408-4400

82:

202-408-4000

TELEPHONE:

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US-08-739C-92
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; Patent No. 6015695...
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FILING DATE: 06 JUNE-1995
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APPLICANT: Casterman, Cecile
APPLICANT: Hamers, Raymond
TITLE OF INVENTION: Immunoglo
NUMBER OF SEQUENCES: 130
CORRESPONDENCE ADDRESS:
  NAME: TRUJILLO, DOREEN YATKO
REGISTRATION NUMBER: 35,719
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LENGTH: 32 amino acids
TYPE: amino acid
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1300 I Street, N.W.
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  95.8%;
93.8%;
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APPLICANT: Bodmer, Mark
APPLICANT: Athwal, Diljeet Singh
APPLICANT: Emtage, John Spencer
APPLICANT: Entage, John Spencer
APPLICANT: TAVENTION: Interleukin-5 Specific Recombinant Antibodies
   PILING DATE: 17-AUG-1993
APPLICATION NUMBER: FR 924
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: FR 934
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PILING DATE: 06-JUN-1995
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   NAME: Potter, Jane REGISTRATION NUMBER:
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Copyright (c) 1993 - 2005 Compugen Ltd.
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| US-10-632-706-196  US-10-632-706-196  Sequence 196, Application US/10632706  Publication No. US20040175385A1  GENERAL INFORMATION:  APPLICANT: MARKS, JAMES D.  APPLICANT: MARKSDORFER, PETER  TITLE OF INVENTION: THERAPBUTIC MONOCLONAL ANTIBODIES THAT NEUTRALIZE BOTULINUM  TITLE OF INVENTION: NEUROTOXINS  FILE REFERENCE: 407T-895120US  CURRENT APPLICATION NUMBER: US/10/632,706  CURRENT APPLICATION NUMBER: US 60/400,721  PRIOR APPLICATION NUMBER: US 60/400,721  PRIOR APPLICATION NUMBER: US 09/144,806 | Query Match 100.0%; Score 165, DB 4; Length 32; Best Local Similarity 100.0%; Pred. No.,3.4e-15; Matches 32; Conservative 0; MismatcHes 0; Indels 0; Gaps 0;  Qy 1 RFTIGRDNSKNTLFLQMNSLRADDTAVYYCAR 32 | US-10-632-706-193 US-10-632-706-193 ; Sequence 193, Application US/10632706 ; Publication No. US20040175385A1 ; GENERAL INFORMATION: APPLICANT: MARKS, JAMES D. APPLICANT: MARKS, JAMES D. TITLE OF INVENTION: THERAPBUTIC MONOCLONAL ANTIBODIES THAT NEUTRALIZE BOTULINUM TITLE OF INVENTION: NEUROTOXINS FILE REFERENCE: 407T-895120US ; CURRENT APPLICATION NUMBER: US/10/632,706 ; CURRENT FILING DATE: 2003-08-01 PRIOR APPLICATION NUMBER: US 60/400,721 ; PRIOR APPLICATION NUMBER: US 60/400,721 ; PRIOR APPLICATION NUMBER: US 99/144,806 ; PRIOR PILING DATE: 1998-08-31 ; NUMBER OF ESQ ID NOS: 278 ; SOFTWARE: Patentin version 3.2 ; LENGTH: 32 ; TYPE: PRT  ORGANISM: Artificial  FEATURE:  OTHER INFORMATION: single chain antibody fragment US-10-632-706-193 | ALIGNMENTS | 28 158 95.8 32 3 US-09-563-242-152 Sequence 152, Appl 158 95.8 32 4 US-10-663-244-28 Sequence 28, Appl 30 158 95.8 32 4 US-10-783-950-142 Sequence 2142, Appl 31 158 95.8 32 5 US-10-784-428-30 Sequence 30, Appl 32 158 95.8 32 5 US-10-751-826-82 Sequence 208, Appl 33 158 95.8 32 5 US-10-728-440B-108 Sequence 108, Appl 34 158 95.8 32 5 US-10-923-068-203 Sequence 203, Appl 35 158 95.8 32 5 US-10-923-068-203 Sequence 204, Appl 36 158 95.8 32 5 US-10-923-068-204 Sequence 204, Appl 37 158 95.8 32 5 US-10-923-068-230 Sequence 204, Appl 38 158 95.8 32 5 US-10-923-068-230 Sequence 335, Appl 38 158 95.8 32 5 US-10-923-068-335 Sequence 335, Appl 39 158 95.8 32 5 US-10-923-068-335 Sequence 336, Appl 39 158 95.8 32 5 US-10-923-068-36 Sequence 356, Appl 39 158 95.8 32 5 US-10-923-068-36 Sequence 362, Appl 39 158 95.8 32 5 US-10-923-068-36 Sequence 362, Appl 39 158 95.8 32 5 US-10-923-068-36 Sequence 362, Appl 39 158 95.8 32 5 US-10-923-068-36 Sequence 362, Appl 39 158 95.8 32 5 US-10-923-068-36 Sequence 362, Appl 39 158 95.8 32 5 US-10-923-068-36 Sequence 362, Appl 39 158 95.8 32 5 US-10-923-068-36 Sequence 362, Appl 39 158 95.8 32 5 US-10-923-068-36 Sequence 362, Appl 39 158 95.8 32 5 US-10-923-068-36 Sequence 362, Appl 30 158 95.8 32 5 US-10-923-068-36 Sequence 362, Appl 30 158 95.8 32 5 US-10-923-068-36 Sequence 362, Appl 30 158 95.8 32 5 US-10-923-068-36 Sequence 362, Appl 30 158 95.8 32 5 US-10-923-068-36 Sequence 362, Appl 30 158 95.8 32 5 US-10-923-068-36 Sequence 362, Appl 30 158 95.8 32 5 US-10-923-068-36 Sequence 362, Appl 30 158 95.8 32 5 US-10-923-068-36 Sequence 362, Appl 30 158 95.8 32 5 US-10-923-068-36 Sequence 362, Appl 30 158 95.8 32 5 US-10-923-068-36 Sequence 362, Appl 30 158 95.8 32 5 US-10-923-068-36 Sequence 362, Appl 30 158 95.8 32 5 US-10-923-068-36 Sequence 362, Appl 30 158 95.8 32 5 US-10-923-068-36 Sequence 362, Appl 30 158 95.8 32 5 US-10-923-068-36 Sequence 362, Appl 30 158 95.8 32 5 US-10-923-068-36 Sequence 362, Appl 30 158 95.8 32 5 US-10-923-068-36 Sequence 362, Appl 30 158 95.8 32 5 US-10-92 |
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TITLE OF INVENTION: THERAPEUTIC MONOCLONAL ANTIBODIES THAT NEUTRALIZE BOTULINUM
FITTLE OF INVENTION: NEUROTOXINS
FILE REFERENCE: 407T-895120US
CURRENT APPLICATION NUMBER: US/10/632,706
CURRENT FILING DATE: 2003-08-01
PRIOR APPLICATION NUMBER: US 60/400,721
PRIOR FILING DATE: 2002-08-01
PRIOR FILING DATE: 2002-08-01
PRIOR FILING DATE: 2002-08-031
NUMBER OF SEQ ID NOS: 278
SOFTWARE: Patentin version 3.2
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Sequence 202, Application US/10632706
Publication No. US20040175385A1
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; OTHER INFORMATION: single chain antibody fragment US-10-632-706-199
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  PRIOR FILING DATE: 1998-08-31
NUMBER OF SEQ ID NOS: 278
SOFTWARE: PatentIn version 3.2
SEQ ID NO 196
LENGTH: 32
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Publication No. US20040175385A1
GENERAL INFORMATION:
APPLICANT: MARKS, JAMES D.
APPLICANT: AMERSDORFER, PETER
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            APPLICANT: MARKS, JAMES D.
APPLICANT: AMERSDORFER, PETER
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TITLE OF INVENTION: NEUROTOXINS
TILE OF INVENTION: NEUROTOXINS
TILE OF INVENTION: NEUROTOXINS
CURRENT APPLICATION NUMBER: US/10/632,706
CURRENT FILING DATE: 2003-08-01
PRIOR APPLICATION NUMBER: US 60/400,721
PRIOR PILING DATE: 2002-08-01
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NUMBER OF SEQ ID NOS: 278
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SOFTWARE: PatentIn version 3.2
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  SEQ ID NO 152
LENGTH: 126
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Publication No. US20030040605A1
GENERAL INFORMATION:
APPLICANT: Slegel, Donald L.
TITLE OF INVENTION: Rh(D)-BINDING PROTEINS AND MAGNETICALLY ACTIVATED CELL
TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF
FILE REFERENCE: 09596-42U2
CURRENT FILING DATE: 2001-05-04
CURRENT FILING DATE: 2001-05-04
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 09/240,274
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 60/028,550
PRIOR FILING DATE: EARLIER FILING DATE: 1996-10-11
NUMBER OF SEQ ID NOS: 224
   Sequence 3, Application US/10886838

Publication No. US20050008642A1

GENERAL INFORMATION:

APPLICANT: HOffmann-La Roche Inc.

TITLE OF INVENTION: Antibodies against insulin-like growth factor I receptor and ITITLE OF INVENTION: thereof

FILE REFERENCE: 21695

CURRENT APPLICATION NUMBER: US/10/886,838

CURRENT FILING DATE: 2003-07-10

RINGER FILING DATE: 2003-07-10

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FEATURE:
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   OTHER INFORMATION: single chain antibody fragment
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96.9%;
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100.0%; Pred. No. 3.4e-15;
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Conservative

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96.98;

Pred. No. 3.9e-14;

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US-09-848-798-139
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  CURRENT APPLICATION NUMBER: US/10/935,290
CURRENT FILING DATE: 2004-09-08
PRIOR APPLICATION NUMBER: PCT/US03/09625
PRIOR FILING DATE: 2003-03-28
PRIOR APPLICATION NUMBER: 60/368,813
PRIOR FILING DATE: 2002-04-01
NUMBER OF SEQ ID NOS: 234
SEQ ID NO 55
   Sequence 55, Application US/10935290
Publication No. US20050069542A1
GENERAL INFORMATION:
APPLICANT: Baker et al.
TITLE OF INVENTION: Antibodies that Specifically Bind to GMAD
FILE REFERENCE: PF584P1
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PRIOR FILING DATE: 2002-12-02
NUMBER OF SEQ ID NOS: 320
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 144
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Best Local Similarity 96.9%;
Matches 31; Conservative
   APPLICANT:
   APPLICANT:
APPLICANT:
APPLICANT:
   APPLICANT: Qiaojuan Jane Su
TITLE OF INVENTION: ANTIBODIES DIRECTED TO TUMOR NECROSIS
TITLE OF INVENTION: FACTOR AND USES THEREOF
FILE REFERENCE: ABGENIX.073A
CURRENT APPLICATION NUMBER: US/10/727,155
CURRENT FILING DATE: 2003.12-02
  APPLICANT: John S. Babcook APPLICANT: Jaspal S. Kang
   APPLICANT:
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  APPLICANT
  PPLICANT
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Xiao Feng
Scott Klakamp
  Craig Pigott
Meina Liang
Rozanne Lee
  Raffaella Faggioni
Giorgio Senaldi
   Orit Foord
   Kathy Manchulencho
  Mary Haak-Frendscho
Palaniswami Rathanaswami
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; Prication Number: US/09/848,798
; CURRENT APPLICATION NUMBER: US/09/848,798
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; PRIOR APPLICATION NUMBER: EARLIER PILING DATE: 1996-10-11
; NUMBER OF ERG ID NOS: 224
; SOFTWARE: Patentin Ver. 2.0

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APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
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APPLICANT:
                               APPLICANT: Kathy Manchulencho
APPLICANT: Raffaella Faggioni
APPLICANT: Giorgio Senaldi
APPLICANT: Giorgio Senaldi
APPLICANT: Qiarginan Jane Su
TITLE OF INVENTION: ANTIBODIES DIRECTED TO TUMOR NECROSIS
TITLE OF INVENTION: PACTOR AND USES THEREOF
FILE REFERENCE: ABGENIX. 073A
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CURRENT FILLING DATE: 2003-12-02
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   APPLICANT:
  APPLICANT:
   John S.
  Xiao Feng
Scott Klakamp
Mary Haak-Frendscho
  Jaspal S. Kang
Orit Foord
  Craig Pigott
Meina Liang
   Palaniswami Rathanaswami
   Application US/10727155

p. US20050049402A1
   Rozanne Lee
   Jarry Green
   Babcook
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APPLICANT: Douglas C. Hanson
APPLICANT: Mark J. Neveu
APPLICANT: Mileen E. Mueller
APPLICANT: Jeffrey H. Hanke
APPLICANT: Jeeven C. Gliman
APPLICANT: C. Geoffrey Davis
APPLICANT: Jose R. Corvalan
APPLICANT: Jose R. Corvalan
   ORGANISM: Homo sapiens -10-727-155-214
   APPLICANT: Raffaella Faggioni
APPLICANT: Glorgio Senaldi
APPLICANT: Olaojuan Jane Su
TITLE OF INVENTION: ANTIBODIES DIRECTED TO TUMOR NECROSIS
TITLE OF INVENTION: FACTOR AND USES THEREOF
FILE REFERENCE: ABGENIX.073A
CURRENT APPLICATION NUMBER: US/10/727,155
CURRENT FILING DATE: 2003-12-02
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   APPLICANT:
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PRIOR FILING DATE: 1998-12-23
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SEQ ID NO 74
  GENERAL INFORMATION:
APPLICANT: Douglas C. Hanson
APPLICANT: Mark J. Neveu
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Matches 31; Conservative
   SOFTWARE: PatentIn Ver. 2.1
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   Sequence 74, Application US/10776649 Publication No. US20040228861A1
APPLICANT: PFIZER PRODUCTS INC.
TITLE OF INVENTION: USES OF ANTI-CTLA-4 ANTIBODIES
FILE REPERENCE: PC23019A
CURRENT APPLICATION NUMBER: US/10/153,382
CURRENT FILING DATE: 2002-05-22
  APPLICANT: C. Geoffrey Davis
APPLICANT: Jose R. COXVALAN
TITLE OP INVENTION: HUMAN MONOCLONAL ANTIBODIES TO CTLA-4
FILE REFERENCE: ABX-PF1 DIV2
CURRENT APPLICATION NUMBER: US/10/776,649
CURRENT FILING DATE: 2004-02-10
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CURRENT FILING DATE: 2003-07-01
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PRIOR FILING DATE: 1999-12-23
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  APPLICANT: Eileen E. Mueller
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APPLICANT: PRIZER PRODUCTS INC.
TITLE OF INVENTION: USES OF ANTI-CTLA-4 ANTIBODIES
FILE REFERENCE: PC23019A
CURRENT APPLICATION NUMBER: US/10/153,382
CURRENT FILING DATE: 2002-05-22
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6

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protein search, using sw model

GenCore version 5.1.6 Copyright (c) 1993 - 2005 Compugen Ltd.

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and is derived by analysis of the total score distribution.
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US-11-177-978-7
US-10-648-816-11
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  7; Search time 0.586081 Seconds (without alignments) 61.686 Million cell updates/sec
   Sequence 10,
Sequence 24,
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sequence
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Sequence
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  Sequence 6,
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  Sequence
6, Appli
10, Appl
24, Appl
16, Appl
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16, Appl
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18, Appl
19, Appl
19, Appl
11, Appl
              APPLICANT: Basi, Guriq
APPLICANT: Saldanha, Jose
APPLICANT: Saldanha, Jose
APPLICANT: Saldanha, Jose
APPLICANT: Yednock, Ted
APPLICANT: Yednock, Ted
TITLE OF INVENTION: HUMANIZED ANTIBODIES THAT RECOGNIZE
TITLE OF INVENTION: BETA-AMYLOID PEPTIDE
FILE REPERENCE: ELAT-002CP
CURRENT APPLICATION NUMBER: US/10/789,273
CURRENT FILING DATE: 2004-02-27
PRIOR APPLICATION NUMBER: US/10/389,389
PRIOR APPLICATION NUMBER: US/10/389,389
PRIOR FILING DATE: 2003-03-12
PRIOR FILING DATE: 2003-03-12
PRIOR FILING DATE: 2003-03-12
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   RESULT 2
US-10-789-273-10
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   Sequence 10, Application US/10789273 Publication No. US20050249725A1
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|------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|-----------------|-----------------|-----------------|-----------------|------------------|------------------|------------------|
|            | 108              | 110              | 110              | 111              | 113              | 114              | 114              | 114              | 114              | 114              | 114              | 118              | 119              | 119             | 126             | 127             | 128             | 128              | 128              | 128              |
|            | 65.5             | 66.7             | 66.7             | 67.3             | 68.5             | 69.1             | 69.1             | 69.1             | 69.1             | 69.1             | 69.1             | 71.5             | 72.1             | 72.1            | 76.4            | 77.0            | 77.6            | 77.6             | 77.6             | 77.6             |
|            | 118              | 313              | 261              | 118              | 119              | 625              | 576              | 328              | 327              | 327              | 258              | 250              | 263              | 119             | 114             | 543             | 121             | 121              | 121              | 118              |
|            | 7                | _                | <b>-</b>         | 7                | _                | μ.               | _                | ۲                | -                | μ.               | μ                | ۲                | -                | 7               | _               | μ               | 7               | μ                | μ                | r                |
| ALIGNMENTS | US-11-012-353-70 | US-10-512-184-72 | US-10-512-184-35 | US-11-012-353-71 | US-10-502-145-25 | US-10-512-184-47 | US-10-512-184-65 | US-10-512-184-63 | US-10-512-184-64 | US-10-512-184-62 | US-10-512-184-26 | US-10-512-184-27 | US-10-512-184-29 | US-11-010-954-5 | US-10-726-554-8 | US-10-495-664-3 | US-11-077-978-3 | US-10-648-816-16 | US-10-648-816-15 | US-10-648-816-14 |
|            | Sequence         | Sequence         |                  | Sequence         | Sequence         | Sequence (       | Sequence         | Sequence         | Sequence (       | Sequence         |                  | Sequence         | Sequence         | Sequence        | Sequence 8      | Sequence        | Seguence        | Sequence 1       |                  | Sequence 1       |
|            | 70,              | 72,              | 35               | 71,              | 25               | 47,              | 65               | 63               | 64               | 62               | 26,              | 27,              | 29,              |                 |                 | ,               |                 | 16,              | , 5              |                  |
|            | Appl             | Appl             | Appl             | Tddv             | Tddv             | Appı             | Appl             | Appı             | Appl             | Tddy             | Tddy             | Appl             | Appl             | Tdd             | tidd            | Tada            | trade           | Tddv             | Appl             | Appl             |

## ; LENGTH: 102 ; TYPE: PRT ; ORGANISM: Homo sapiens US-10-997-201A-6 US-10-997-201A-6 APPLICANT: Maragoo, Wayne APPLICANT: Sui, Jianhua TITLE OF INVENTION: Antibodies Against SARS-COV and Methods of Use Thereof FILE REFERENCE: 20363-026 CURRENT APPLICATION NUMBER: US/10/997,201A CURRENT FILING DATE: 2004-11-24 PRIOR APPLICATION NUMBER: 60/524,840 PRIOR APPLICATION NUMBER: 60/524,840 PRIOR FILING DATE: 2003-11-25 NUMBER OF SEQ ID NOS: 40 SOFTWARE: Patentin Ver. 2.1 Sequence 6, Application US/10997201A Publication No. US20050249739A1 GENERAL INFORMATION: Query Match Best Local Matches 57 30; Similarity Conservative 95.8%; Score 158; DB 1; 1 Pred. No. 9.2e-13; 2; Mismatches 0; Length 102; Indels 0 Gaps 0

PRIOR APPLICATION NUMBER: US 60/251,892

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APPLICANT: Cohen, Bruce D.

APPLICANT: Beabe, Jean

APPLICANT: Miller, Penelope B.

APPLICANT: Moyer, James D.

APPLICANT: Moyer, James D.

APPLICANT: Corvalan, Jose R.

APPLICANT: Corvalan, Jose R.

APPLICANT: Corvalan, Jose R.

APPLICANT: Corvalan, Jose R.

APPLICANT: ANTIBODIES TO INSULIN-LIKE GROWTH FACTOR I RECEPTOR FILE REFERENCE: ABX-PF2

CURRENT APPLICATION NUMBER: US/11/144,248

CURRENT APPLICATION NUMBER: US/10/038,591

PRIOR APPLICATION NUMBER: US/10/038,591

PRIOR APPLICATION NUMBER: 60/259,927

PRIOR FILING DATE: 2002-01-05
  ; PRIOR FILING DATE: 2000-12-06; NUMBER OF SEQ ID NOS: 63; SOFTWARE: FastSEQ for Windows Version 4.0; SEQ ID NO 10; SEQ ID NO 10; LENGTH: 98; TYPE: PRT; ORGANISM: Homo sapiens
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   US-11-144-248-24
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   CURRENT APPLICATION NUMBER: US/11/144,248
CURRENT FILING DATE: 2005-06-02
PRIOR APPLICATION NUMBER: US/10/038,591
PRIOR FILING DATE: 2002-01-04
PRIOR APPLICATION NUMBER: 60/259,927
PRIOR PILING DATE: 2001-01-05
NUMBER: 2001-01-05
   APPLICANT: Cohen, Bruce D.
APPLICANT: Beebe, Jean
APPLICANT: Miller, Penelope B.
APPLICANT: MOYEr, James D.
APPLICANT: Corvalan, Jose R.
APPLICANT: Gallo, Michael
APPLICANT: Gallo, Michael
APPLICANT: MILENTION: ANTIBODIES TO INSULIN-LIKE GROWTH FACTOR I RECEPTOR
   FILE REFERENCE: ABX-PF2
   ORGANISM: Homo sapiens
  LENGTH: 98
  Local Similarity
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90.6%; Pred. No. 1.9e-12;
   93.9%; Score 155; DB 1; Length 98; 90.6%; Pred. No. 1.9e-12; tive 3; Mismatches 0; Indels
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US-10-721-763-33

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TITLE OF INVENTION: ANTI TRAIL-R ANTIBODY
FILE REFERENCE: PH-1573-PCT
CURRENT APPLICATION NUMBER: US/10/721,763
CURRENT FILING DATE: 2003-11-26
PRIOR APPLICATION NUMBER: JP2001-150213
PRIOR FILING DATE: 2001-05-18
PRIOR APPLICATION NUMBER: JP2001-243040
PRIOR APPLICATION NUMBER: JP2001-243040
PRIOR APPLICATION NUMBER: JP2001-314489
PRIOR APPLICATION NUMBER: JP2001-314489
PRIOR FILING DATE: 2001-10-11
NUMBER OF SEQ ID NOS: 45
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; ORGANISM: Homo sapiens
US-11-144-248-16
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US-11-144-248-16
   US-11-144-248-24
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   APPLICANT: Corvalan, Jose R.
APPLICANT: Gallo, Michael
TITLE OF INVENTION: ANTIBODIES TO INSULIN-LIKE GROWTH FACTOR I RECEPTOR
FILE REFERENCE: ABX-PF2
CURRENT APPLICATION NUMBER: US/11/144,248
CURRENT FILING DATE: 2005-06-02
PRIOR APPLICATION NUMBER: US/10/038,591
PRIOR APPLICATION NUMBER: US/10/038,591
PRIOR APPLICATION NUMBER: 60/259,927
PRIOR APPLICATION NUMBER: 60/259,927
PRIOR FILING DATE: 2001-01-05
PRIOR FILING DATE: 2001-01-05
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  NUMBER OF SEQ ID NOS: 60
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   APPLICANT: Cohen, Bruce D.
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Miller, Penelope B.
Moyer, James D.
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   93.9%; Score 155; DB 7; Length 125; 90.6%; Pred. No. 2.4e-12;
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Pred. No. 2.4e-12;
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; TYPE: PRT ; ORGANISM: Homo sapiens US-10-721-763-33

Query Match Best Local Matches 2

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Conservative

Mismatches

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Local Similarity

93.9%;

Score 155; DB 1; Length 139; pred. No. 2.7e-12;

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CURRENT APPLICATION NUMBER: US/11/144,248
CURRENT FILING DATE: 2005-06-02
PRIOR APPLICATION NUMBER: US/10/038,591
PRIOR FILING DATE: 2002-01-04
PRIOR APPLICATION NUMBER: 60/259,927
PRIOR PILING DATE: 2001-01-05
NUMBER OF SEQ ID NOS: 60
SOFTWARE: PALENTIN Ver. 2.1
SEQ ID NO 46
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  TITLE OF INVENTION: HUMANIZED ANTIBODIES THAT RECOGNIZE
TITLE OF INVENTION: BETA-ANYLOID PEPTIDE
FILE REFERENCE: ELN-002CP
CURRENT APPLICATION NUMBER: US/10/789,273
CURRENT FILING DATE: 2004-02-27
PRIOR APPLICATION NUMBER: US/10/389,389
PRIOR FILING DATE: 2003-03-12
PRIOR PILING DATE: 2003-03-12
PRIOR APPLICATION NUMBER: US/10/10/942
PRIOR APPLICATION NUMBER: US/10/10/942
PRIOR FILING DATE: 2001-12-06
PRIOR FILING DATE: 2001-12-06
PRIOR FILING DATE: 2000-12-06
   Sequence 46, Application US/11144248 Publication No. US20050244408A1
NUMBER OF SEQ ID NOS: 63
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 12
LENGTH: 138
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  -11-144-248-46
  GENERAL INFORMATION:
APPLICANT: Basi, Guriq
APPLICANT: Saldanha, Jos
APPLICANT: Yednock, Ted
  -11-144-248-46
  APPLICANT: Cohen, Bruce D.
APPLICANT: Beebe, Jean
APPLICANT: Miller, Penelo
  RPPLICANT: Gallo, MICHAEL
TITLE OF INVENTION: ANTIBODIES TO INSULIN-LIKE GROWTH FACTOR I RECEPTOR
  TYPE: PRT
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Miller, Penelope B.
Moyer, James D.
  Corvalan, Jose R. Gallo, Michael
  RFTISRDNSKNTLYLQMNSLRAEDTAVYYCAK 117
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US-10-789-273-12
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US-11-144-248-30
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  US-11-144-248-30
  APPLICANT: Adolf, Guenther
APPLICANT: Heider, Karl-Heinz
APPLICANT: Heider, Karl-Heinz
APPLICANT: Heider, Karl-Heinz
APPLICANT: Heider, Karl-Heinz
APPLICANTON: Compositions and Methods fo
TITLE OF INVENTION: Cytotoxic CD44 Antibody Im
TITLE OF INVENTION: Cytotoxic CD44 Antibody Im
TITLE OF INVENTION: Cytotoxic CD44 Antibody Im
TITLE OF INVENTION: Cytotoxic CD44 Antibody Im
TITLE OF INVENTION: Cytotoxic CD44 Antibody Im
TITLE OF INVENTION: Cytotoxic CD44 Antibody Im
PRICE REPERENCE: 1/1383
CURRENT APPLICATION NUMBER: US/11/172,320
CURRENT FILING DATE: 2003-08-21
PRIOR APPLICATION NUMBER: US/10/645,215
PRIOR APPLICATION NUMBER: ED 02 018 686.2
   US-11-172-320-6
   FILE REFERENCE: ABX-PF2
CURRENT APPLICATION NUMBER: US/11/144,248
CURRENT FILING DATE: 2005-06-02
PRIOR APPLICATION NUMBER: US/10/038,591
PRIOR FILING DATE: 2002-01-04
PRIOR APPLICATION NUMBER: 60/259,927
PRIOR FILING DATE: 2001-01-05
PRIOR FILING DATE: 2001-01-05
NUMBER OF SEQ ID NOS: 60
SOFTWARE: Patentin Ver. 2.1
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   Sequence 6, Application US/11172320 Publication No. US20050244413A1
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  APPLICANT:
  APPLICANT: Miller, Penelope E.
APPLICANT: Moyer, James D.
APPLICANT: Corvalan, Jose R.
APPLICANT: Gallo, Michael
TITLE OF INVENTION: ANTIBODIES TO INSULIN-LIKE GROWTH FACTOR I RECEPTOR
   APPLICANT: Cohen, Bruce D.
   FEATURE:
   ORGANISM: Artificial Sequence
  OTHER INFORMATION: Humanized 3D6 light chain variable region
APPLICATION NUMBER: EP 02 018 686.2 FILING DATE: August 21, 2002 APPLICATION NUMBER: US 60/405,956
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  US-11-144-248-50
  Sequence 4, Application US/11144248
Publication No. US20050244408A1
GENERAL INFORMATION:
APPLICANT: Beebe, Bruce D.
APPLICANT: Beebe, Jean
APPLICANT: Miller, Penelope E.
APPLICANT: Moyer, James D.
APPLICANT: Corvalan, Jose R.
APPLICANT: Corvalan, Jose R.
APPLICANT: Gallo, Michael
TITLE OF INVENTION: ANTIBODIES TO INSULIN-LIKE GROWTH FACTOR I RECEPTOR
FILE REFERENCE: ABS-PF2
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CURRENT APPLICATION NUMBER: US/11/144,248 CURRENT FILING DATE: 2005-06-02 PRIOR APPLICATION NUMBER: US/10/038,591 PRIOR FILING DATE: 2002-01-04 PRIOR APPLICATION NUMBER: 60/259,927
  TITLE OF INVENTION: ANTIBODIES TO INSULIN-LIKE GROWTH FACTOR I RECEPTOR FILE REFERENCE; ABX-PF2 CURRENT APPLICATION NUMBER: US/11/144,248 CURRENT FILING DATE: 2005-06-02 PRIOR APPLICATION NUMBER: US/10/038,591 PRIOR FILING DATE: 2002-01-04 PRIOR FILING DATE: 2002-01-04
   APPLICANT: Cohen, Bruce D.
APPLICANT: Beebe, Jean
APPLICANT: Miller, Penelop
APPLICANT: Moyer, James D.
  PRIOR FILING DATE: 2001-01-05
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   APPLICANT: Corvalan, Jose R. APPLICANT: Gallo, Michael
   TYPE: PRT
ORGANISM: Artificial Sequence
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Similarity 87.5%;
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  RESULT 13
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APPLICANT: Beebe, Jean
APPLICANT: Miller, Penelope E.
APPLICANT: Miller, Penelope E.
APPLICANT: Moyer, James D.
APPLICANT: Gorvalan, Jose R.
APPLICANT: Gallo, Michael
TITLE OF INVENTION: ANTIBODIES TO INSULIN-LIKE GROWTH FACTOR I RECEPTOR
EILE REFERENCE: ABX-PF2
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CURRENT FILING DATE: 205-06-02
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CURRENT FILING DATE: 2005-06-02
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PRIOR PILING DATE: 2002-01-04
PRIOR PILING DATE: 2002-01-04
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NUMBER OF SEQ ID NOS: 60
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  APPLICANT: Miller, Penelope E.
APPLICANT: Moyer, James D.
APPLICANT: Corvalan, Jose R.
APPLICANT: Gallo, Michael
   APPLICANT: Cohen, Bruce D. APPLICANT: Beebe, Jean
PRIOR APPLICATION NUMBER: US/10/038,591
PRIOR FILING DATE: 2002-01-04
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PRIOR FILING DATE: 2001-01-05
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GENERAL INFORMATION:
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APPLICANT: Beebe, Jean
APPLICANT: Miler, Penelope E.
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APPLICANT: Moyer, James D.
APPLICANT: Gallo, Michael
ITITLE OF INVENTION: ANTIBODIES TO INSULIN-LIKE GROWTH FACTOR I RECEPTOR
FILE REFERENCE: ABX-PF2
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CURRENT FILING DATE: 2005-06-02
PRIOR APPLICATION NUMBER: US/10/038,591
PRIOR FILING DATE: 2001-01-04
PRIOR APPLICATION NUMBER: 60/259,927
PRIOR APPLICATION NUMBER: 60/259,927
PRIOR RILING DATE: 2001-01-05
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